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; Sequence 2zu, ...
; Patent No. 6262334
; Patent No. 6262334
; Patent No. 6262334
; GENERAL INFORMATION:
; APPLICANT: ENDEGE, WILSON O., ET AL.
; TITLE OF INVENTION: NOVEL HUMAN GENES AND TITLE OF INVENTION: PRODUCTS: II
; FILE REFERENCE: CCDNA-260XX
CURRENT APPLICATION NUMBER: US/09/385,9
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APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUC
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
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Best Local S
Matches 32
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Best Local
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ORGANISM: Homo s
-09-182-145-14
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SEQ ID NO 14
LENGTH: 1293
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NAME/KEY:
LOCATION:
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Hillan, Kennetl
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Levine, Arnold J.
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e, David A.
                                       NUMBER: US/09/385,982
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APPLICANT: Wood, William I.
FITLE OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC AC
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-02-04
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
                                                                           ORGAN-
FEATURE:
NAME/KEY: misc____L
LOCATION: 1-841
OTHER INFORMATION: $
~nt No. 6387657
                                                                                             Patent No
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INFORMATION: n
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APPLICATION NUMBER: 09/328,111
FILING DATE: 1999-06-08
APPLICATION NUMBER: 60/117,393
                                                                                                                                                                                                                                                                                                                                                                                                    Cohen, KODELL
Goddard, Audrey
Gurney, Austin L.
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9; Conservative
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misc_
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1-841
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imilarity 100.0%;
Conservative
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Lawrence, David A.
Levine, Arnold J.
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RESULT 9

US-09-149-46-225/c

US-09-149-476-225/c

US-09-149-476-225/c

GENERAL INFORMATION: APPLICANTON: 186 Human Secreted proteins FILE REPERENCE: P2002P1

CURRENT HAPLICATION NUMBER: US/09/149,476

CURRENT HILING DATE: 1999-03-06

EARLIER APPLICATION NUMBER: 60/040,162

EARLIER FILING DATE: 1997-03-07

EARLIER PELING DATE: 1997-03-07

EARLIER FILING DATE: 1997-05-23

EARLIER FILING DATE: 1

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1997-08-22 NUMBER: 60/047,59	1997-08-22	: 1997-08-22 NIMBER: 60/056.892	BER: 60	NUMBER: 60	NUMBER: 60		7-08-22	UMBER: 60/056,63 1997-08-22	1997-08-22	997-08-22 MBER: 60/056	MBER: 60	NUMBER: 60,	NUMBER: 60,	: 1997-08-22	1997-08-22	997-08-22	: 1997-08-22 NUMBER: 60/056,637	BER: 60	NUMBER: 60	NUMBER: 60, 1997-08-2	NUMBER: 60/056,878: 1997-08-22	997-08-22	997-08-22	997-08-22	997-08-22 MBER: 60/056,	997-08- MBER: 6	NUMBER: 6	NUMBER: 60	NUMBER	NUMBER: 6	NUMBER	NUMBER: 60/043,312 : 1997-04-11	NUMBER: 60/043,669: 1997-04-11	997-04-11	997-04-11	: 1997-04-11 NUMBER: 60/043,671	: 199/-04-11 NUMBER: 60/043,311	NUMBER	NUMBER	NUMBER: 60/043,568: 1997-04-11

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RESULT 10
US-09-149
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RESULT 10

JS-09-149-476-57/c

Sequence 57, Application (
Patent No. 6420526

GENERAL INFORMATION:

APPLICANT: Rosen et al.
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Matches
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REFILING DATE: 1997-04-15
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REFILING DATE: 1997-04-16
REFILING DATE: 1997-04-17
REFILING DATE: 1997-04-22
REFILING DATE: 1997-08-22
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FILE REFERENCE: PEOD2P1
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT APPLICATION NUMBER: US/09/149,449
BARLIER FILING DATE: 1999-03-06
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,33
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-03-23
EARLIER FILING DATE: 1997-0
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FILING DATE: 1997-04
APPLICATION NUMBER: 1997-04
APPLICATION NUMBER: FILING DATE: 1997-04
APPLICATION NUMBER: APPLICATION NUMBER: 1997-04

. 1997 NUMBER: 1997-0

-04-11 R: 60/ -04-11

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R APPLICATION NUMBER: 60/043, 31
R FILING DATE: 1997-04-11
R APPLICATION NUMBER: 60/043, 31
R FILING DATE: 1997-06-06
R APPLICATION NUMBER: 60/056, 88
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 88
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R APPLICATION NUMBER: 60/056, 88
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R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 68
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 68
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 88
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 88
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 89
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R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056, 89
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/056
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JS-09-213-768-1
Sequence 1, A
Patent No. 59
GENERAL INFOR
APPLICANT: B
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Best Loo
Matches
      APPLICANT:
TITLE OF IN
FILE REFERE
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FIL:
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      FORMALL...
: Brenda F. Baker
: Lex M. Cowsert
: Lex M. Cowsert
INVENTION: ANTISENSE MODULATION OF SIRENCE: RTS-0026
APPLICATION NUMBER: US/09/213,768
TITTING DATE: 1998-12-17
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PPLICATION NUMBER: 60/056,664

ILING DATE: 1997-08-22

PPLICATION NUMBER: 60/056,876

ILING DATE: 1997-08-22

PPLICATION NUMBER: 60/056,881

ILING DATE: 1997-08-22

PPLICATION NUMBER: 60/056,909

ILING DATE: 1997-08-22

PPLICATION NUMBER: 60/056,875

ILING DATE: 1997-08-22

PPLICATION NUMBER: 60/056,862

ILING DATE: 1997-08-22

ILING DATE: 1997-08-22
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PLICATION NUMBER: 60/049,610
PLICATION NUMBER: 60/049,610
PLING DATE: 1997-06-13
PLING DATE: 1997-06-13
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PPLICATION NUMBER: 60/047,614

ILING DATE: 1997-05-23

PPLICATION NUMBER: 60/043,578

ILING DATE: 1997-04-11

PPLICATION NUMBER: 60/043,576

ILING DATE: 1997-04-11

PPLICATION NUMBER: 60/047,501

ILING DATE: 1997-05-23

PPLICATION NUMBER: 60/043,670

ILING DATE: 1997-04-11

PPLICATION NUMBER: 60/043,670

ILING DATE: 1997-04-11

ILING DATE: 1997-04-11

ILING DATE: 1997-04-11

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ILING DATE: 1997-04-11
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PLICATION NUMBER: 60/0
LING DATE: 1997-09-05
PLICATION NUMBER: 60/0
LING DATE: 1997-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ING DATE: 1997-08-22
LICATION NUMBER: 60/056,908
ING DATE: 1997-08-22
LICATION NUMBER: 60/048,964
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ING DATE: 1997-05-23
LICATION NUMBER: 60/047,589
ING DATE: 1997-05-23
LICATION NUMBER: 60/047,593
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985664
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LICATION NUMBER:
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1997-05-23
NUMBER: 60/047,614
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Pred. No.
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                                   ; Sequence 2, Application US/09213; Patent No. 5985664; GENERAL INFORMATION:
APPLICANT: Brenda F. Baker; APPLICANT: Lex M. Cowsert; TITLE OF INVENTION: ANTISENSE MITTLE REFERENCE: RTS-0026
                                                                                                                                                             В
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; LOCATION: (136)..
US-09-213-768-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wang, Jian-Rui
APPLICANT: Xue, Aidong J.
APPLICANT: Xu, Chongjun
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: No. 6436703el Nucleic &
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 790CIP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT FILING DATE: 2000-09-22
                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                      Best Loc
Matches
                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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Best Local
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      CURRENT APPLICATION NUMBER: US/09/213,768
CURRENT FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 47
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                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ
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PRIOR FILING DATE: 2000-03-31
                                                                                                                                                                                                                                                NAME/KEY: CDS
LOCATION: (13
09-668-680-13
                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: FEATURE:
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ORGANISM:
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Similarity 100.0%;
19; Conservative
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19; Conservative
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Liu, Chenghua
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Pred. No. 4.1
0; Mismatches
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                                                   MODULATION
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Pred. No
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4.1;
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                                                   EXPRESSION
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RESULT
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OTHER INFORMATION: PCR Primer US-09-213-768-2
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Best Local :
                                                      INFORMATION
SEQUENCE
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APPLICANT:
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APPLICANT:
APPLICANT:
                                               APPLICATION NUMBER: US 08/086,415
FILING DATE: 01-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: Daniel W. Collins
REGISTRATION NUMBER: 31,912
REFERENCE/DOCKET NUMBER: 5381.US.P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 937-6365
TELEPHONE: (708) 938-2623
IFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                           TELECOMMU
TELEPHO
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ATTORNEY/
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APPLICANT:
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                                                                                                                                                                           CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: One Abbott Park Road
CITY: Abbott Park
STATE: Illinois
                                                                                                                                                                                                                                       CURRENT
                                                                                                                                                                                                                                                                                                               COMPUTER
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                               TYPE
                                                                                                                                                                                                              SOFTWARE: WordPerfect 5.1
JRRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
FILING DATE: 20-JUN-1994
                                                                                                                                                                                                                                                                                                                            ZIP:
                                                                                                                                                                                                                                                               OPERATING SYSTEM:
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                                            LENGTH
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                                                                                                                                                                                                                                                                                                    MEDIUM
         TOPOLOGY:
                     STRAND
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AL INFORMATION:
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Motley, Cheryl T.
Peterson, Bryan
Edwards, Michelle
Guidinger, Peggy
Tate, Cynthia
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Rosenblatt, Joseph D.
Chen, Irvin S. Y.
Golde, David W.
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Sequence ...

Patent No. 5827685

GENERAL INFORMATION:

APPLICANT: Lindquist, Susan

TITLE OF INVENTION: Methods and Compositions of.

TITLE OF INVENTION: Stress Response Systems

NUMBER OF SEQUENCES: 2

CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold, White & Durkee

STREET: P. O. Box 4433

CITY: Houston

STATE: TX

COUNTRY: USA

ZIP: 77210

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWAREE: Patentin Release #1.0, Version #1.

CURRENT APPLICATION. NUMBER: US/08/249,380

FILING DATE: 31-MAY-1991

ATTORNEY/AGENT INFORMATION:

APPLICATION NUMBER: US/08/249,380

FILING DATE: 31-MAY-1991

ATTORNEY/AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: ARCD:024

TELEPHONE: 713-787-1400

TELEFAX: 713-789-2679

INFORMATION OF SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:

LENGTH: 3727 base pairs

TYPE: nucleic acid

STRANDEDNESS: Single

TOPOLOGY: linear

MOLECULE TYPE: DNA

US-08-249-380-1

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100.08; Pred. NO. 13;
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US-08-249-380-1/c
; Sequence 1, Application US/08249380
; Patent No. 5827685
; GENERAL INFORMATION:
APPLICANT: Lindquist, Susan
Search completed: July Job time: 35.7003 secs
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Best Local Similarity 100.0%;
Matches 18; Conservative
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Published_Applications_NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

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11: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

15: /cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*

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## SUMMARIES

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SEQUENCE CHARACTERISTICS:	INFORMATION FOR SEQ ID NO: 1:	TELEFAX: (617)742-4214	TELEPHONE: (617)227-7400	TELECOMMUNICATION INFORMATION:			NAME: ANY E. MANUTANCO SC 207	NAME. AND EL MOSCHOOLSON	ATTORNEY /AGENT INFORMATION:		APPLICATION NUMBER: <unknown></unknown>	FILING DATE: March 19, 1998	APPLICATION NUMBER: 09/044,273		CLASSIFICATION: <unknown></unknown>	FILING DATE: 07-Dec-2001	APPLICATION NUMBER: US/10/010,408		SUPERMARE: PACEFICIT RELEASE #1.0, VERSION #1.20	Datorem. rc-boo/mo-boo	Z.		COMPUTER READABLE FORM:	ZIP: 02109	COUNTRY: USA		CITY: Boston	State Str	ADDRESSEE: LAHIVE & COCKFIELD, LLP	$\Box$	NUMBER OF SEQUENCES: 13		: No. US2002	APPLICANT: John J. Castellot, Jr.	GENERAL INFORMATION:	Sequence 1, Application US/10010408	US-10-010-408-1	RESULT 1	

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RESULT 2
US-10-010-408-3
; Sequence 3, Application US/10010408
; Publication No. US20020165185A1
; GENERAL INFORMATION:
APPLICANT: John J. Castellot,
TITLE OF INVENTION: No. US2002
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NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & C
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CITY: Boot
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TYPE: nucleic acic
STRANDEDNESS: sinc
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FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
CRAPLICATION ATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
CRNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
ECOMMUNICATION INFORMATION:
TELEPHONE: (617)742-4214
ION FOR SEQ ID NO: 3:
UENCE CHARACTERISTICS:
LENGTH: 753 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
ECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
LOCATION:
                                                  GAGGTGAATGGCCGCAGGTACCTGGATGGAGAGACCTTTAAACCCAATTGCAGGGTCCTG
                                                                                                        CAGGGGGTACCCCTGGTGCTGGATGGCTGTGGCTGCTGTAAAGTGTGTGCACGGAGGCTG
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS
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NT APPLICATION DATA:
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ZIP: 02109
TGCCGCTGTGATGACGGTGGCTTCACC
                          TGCCGCTGTGATGACGGTGGCTTCACC 635
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illarity 100.0%;
Conservative 0
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Pred. No. 2.6e-194;
0; Mismatches 0;
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12 Application

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Best Local S
Matches 318
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                                                                                                                                                                                                                                                                                                                      LOCATION: 1..681
SEQUENCE DESCRIPTION: 10-010-408-12
                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617)742-42
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 681 base pai
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
FORMATION SEQ ID NO: 12:
SECURENCE CHARACTERISCE.
                                                                                                                                                                                                                                                                               y Match
Local Similarity 100.0%;
hes 318; Conservative
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COMPUTER READABLE E
MEDIUM TYPE: E
COMPUTER: IBM
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CORRESPONDENCE
ADDRESSEE
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: cDNA
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CITY: Boston
STATE: Massac
COUNTRY: USA
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GATGACGGTGGCTTCACC
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                                    INVENTION:
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NVENTION: No. US20020165185Alel
and Uses Therefor
SEQUENCES: 13
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Y: USA
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PE: Floppy of IBM PC com
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linear
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7.4e-158;
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RESULT 4
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US-10-010-408
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Best Local S
Matches 210
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APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 8:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
RRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                             NAME/KEY: CDS
LOCATION: 1..210
UENCE DESCRIPTION: 98-8
                                                                                                                                                                                                          Similarity Conserv
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JTER READABLE FORM:

MEDIUM TYPE: Flopp
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STATE: Massachusetts
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ESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10010408
No. US20020165185A1
           CCTGGCGGCCATGGGGCTGTGTGTCTCTTG
                                                  CCCCTGGTGCTGGATGGCTGTGGCTGCTGTAAAGTGTGTGCACGGAGGCTGGGGGAGTCC
                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
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CCTGGCGGCCATGGGGCTGTGTGTCTCTTG
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OF INVENTION: No. US20020165185Alel Heparin-Induced CCN-Like
and Uses Therefor
OF SEQUENCES: 13
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ilarity 100.0%;
Conservative 0
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US-10-010-408-
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                                                 Sequence 17, Applica Publication No. US20 GENERAL INFORMATION:
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-00/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)742-4214
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                Query Match
Best Local
                         APPLICANT:
                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: 1..177
SEQUENCE DESCRIPTION:
10-010-408-5
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: John J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE
                                                                                                                                                                                                                                                                                                                                                            FEATURE
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                                                                                                                                                                        909
                                                                                                                                                                                                                            546
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                         Botstein,
Cohen, R
                                                                                                                                            CTGTGCCGCTGTGATGACGGTGGCTTCACC
                                                                                                                                                                                                LENGTH: 177 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/10010408 No. US20020165185A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS
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Goddard,
Gurney, /
                                                              Application US/10112267
No. US20030068678A1
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                          Robert
Audrey
Austin I
                                      David
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: No. US20020165185A1el
and Uses Therefor
                                                                                                                                                                                                                                                                                                                      SEQ ID
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Pred. No
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                                                                                                                                                                                                                                                                  NO.
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FILE REFERENCE: P1176R2

CURRENT APPLICATION NUMBER: US/10/112,267

CURRENT FILING DATE: 2002-03-27

PRIOR APPLICATION NUMBER: EARLIER APPLICATION

PRIOR APPLICATION NUMBER: EARLIER APPLICATION

PRIOR FILING DATE: EARLIER FILING DATE: 1997

PRIOR APPLICATION NUMBER: EARLIER APPLICATION

PRIOR APPLICATION NUMBER: EARLIER APPLICATION

PRIOR FILING DATE: EARLIER FILING DATE: 1997

PRIOR APPLICATION NUMBER: EARLIER APPLICATION

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PRIOR FILING DATE: EARLIER FILING DATE: 1997

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PRIOR FILING DATE: 1997
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US-10-1
US-10-112-267
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CURRENT FILING DATE: 2002-03-27

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US

PRIOR FILING DATE: EARLIER FILING DATE: 1998-10-29

PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-29

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US

PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-04

PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-14

NUMBER OF SEQ ID NOS: 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM:
US-10-112-267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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Best Local
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                                     NUMBER OF SE
SEQ ID NO 18
LENGTH: 173
TYPE: DNA
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ICANT: Wood, William I.
E OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC
REFERENCE: P1176R2
                                                                                                                                                                                                                                                                                                                                        CANT: Roy, Margaret Ann
CANT: Wood, William I.
OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC ACIDS
REFERENCE: P1176R2
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Pennica, Diane
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                     musculus
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, David A.
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Pred. No. 2e-37;
0; Mismatches
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APPLICATION NUMBER: IDATE: 1997-10-29
                                                                                                                            APPLICATION NUMBER:
DATE: 1998-04-14
                                                                                                                                                                    DATE: 1998-02-04
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CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION
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PRIOR FILING DATE: EARLIER FILING DATE: 1997
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION
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NUMBER OF SEQ ID NOS: 156
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Best Local S
Matches 32
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APPLICANT: Baker, Ke
APPLICANT: Beresin
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TYPE: DNA
ORGANISM: Homo
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Desnoyers, Luc
Filvaroff, Ellen
Gao, Wei-Qiang
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A
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Goddard,
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Hillan, Kenneth J.
Lawrence, David A.
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R FILING DATE: 1997-09-18
R APPLICATION NUMBER: 60/059352
R FILING DATE: 1997-09-19
R APPLICATION NUMBER: 60/059352
R FILING DATE: 1997-09-24
R APPLICATION NUMBER: 60/06250
R FILING DATE: 1997-10-17
R APPLICATION NUMBER: 60/06285
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R APPLICATION NUMBER: 60/063550
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R APPLICATION NUMBER: 60/063561
R APPLICATION NUMBER: 60/063704
R APPLICATION NUMBER: 60/063704
R APPLICATION NUMBER: 60/063733
R FILING DATE: 1997-10-29
R APPLICATION NUMBER: 60/063735
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R APPLICATION NUMBER: 60/056974
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R FILING DATE: 1997-08-26
R FILING DATE: 1997-09-17
R APPLICATION NUMBER: 60/059115
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R APPLICATION NUMBER: 60/059117
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NVENTION: SECRETED AND TRANSMEMBRANE
NVENTION: ACIDS ENCODING THE SAME
                                                                              WG DATE: 1997-10-17
[CATION NUMBER: 60/064248
WG DATE: 1997-11-03
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CATION NUMBER: 60/056974
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Wood,William
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NUMBER: 60/C
                                                      TUMBER: 60/064809
1997-11-07
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## FILING DATE: 1997-11-24
## APPLICATION NUMBER: 60,0692
## FILING DATE: 1997-12-11
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S-10-146-726-3
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R APPLICATION NUMBER: 60/086430

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R APPLICATION NUMBER: 60/087106

R FILING DATE: 1998-05-28

R APPLICATION NUMBER: 60/088026

R FILING DATE: 1998-06-04

R APPLICATION NUMBER: 60/088730

R FILING DATE: 1998-06-10

R APPLICATION NUMBER: 60/088810

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R APPLICATION NUMBER: 60/088858

R FILING DATE: 1998-06-11

R APPLICATION NUMBER: 60/089532

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R APPLICATION NUMBER: 60/089599

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R APPLICATION NUMBER: 60/089907

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R APPLICATION NUMBER: 60/089947

PR FILING DATE: 1998-06-24

PR APPLICATION NUMBER: 60/090429

R FILING DATE: 1998-06-24

R APPLICATION NUMBER: 60/09045

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Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
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Beresini, Maureen
DeForge, Laura
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imilarity 100.0%;
Conservative
                               Zhang, Zemin
VENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
VENTION: ACIDS ENCODING THE SAME
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Wood, William
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Filvaroff, Ellen
Gao, Wei-Qiang
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D. US20030129690A1
   ON: ACIDS
P3330R1C308
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9.2e-07;
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RESULT 11
US-10-146-727-319
; Sequence 319, Application US/10146727
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                                                                                RESULT 12
US-10-146-788-319
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                                                 Sequence 319, Application US/10146788 Publication No. US20030129693A1 GENERAL INFORMATION:
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CURRENT FILING DATE:
Prior Application rem
NUMBER OF SEQ ID NOS:
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Best Local
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ORGANISM: HOMO
-10-146-727-319
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TYPE: DNA
ORGANISM: Homo 9
-10-146-726-319
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Prior Application removed - See
NUMBER OF SEQ ID NOS: 550
                                                                                                                                                                                                                                                                                                           APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED
TITLE OF INVENTION: ACIDS EN
FILE REFERENCE: P3330R1C312
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Q ID NO 319
LENGTH: 1266
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Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy
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Wood, William
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Filvaroff, Ellen
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; TYPE: DNA; ORGANISM: US-10-152-380-
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US-10-152-380
                                                                                                  FILE OF INVENTION.
FILE REFERENCE: P3330R1C397
CURRENT APPLICATION NUMBER: US/10/152,380
CURRENT FILING DATE: 2002-05-21
Prior Application removed - See Palm or File
NUMBER OF SEQ ID NOS: 550
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TYPE: DNA
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NT: Filvaroff, Ellen
NT: Gao, Wei-Qiang
NT: Gao, Wei-Qiang
NT: Goddard, Audrey
NT: Goddard, Audrey
NT: Goddard, Austin L.
NT: Sherwood, Steven
NT: Sherwood, Steven
NT: Stewart, Timothy A.
NT: Stewart, Timothy A.
NT: Tumas, Daniel
NT: Watanabe, Colin K
NT: Watanabe, Colin K
NT: Wood, William
NT: Zhang, Zemin
NT: Zhang, Zemin
SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FINVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FINVENTION: ACIDS ENCODING THE SAME
SEFERCE: P333OR1C397
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE PO
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C412
CURRENT APPLICATION NUMBER: US/10/153,934
CURRENT APPLICATION NUMBER: US/10/153,934
CURRENT FILING DATE: 2002-05-22
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NUMBER OF SEQ ID NOS: 550
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LENGTH: 1266
TYPE: DNA
ORGANISM: Homo Sapien
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## ALIGNMENTS

FEATURES	COMMENT	JOURNAL	TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	Locus	BQ93/88/
Tissue Procurement: The Cepko Laboratory  cDNA Library Preparation: Life Technologies, Inc.  cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)  DNA Sequencing by: Agencourt Bioscience Corporation  Clone distribution: MGC clone distribution information can be  found through the I.M.A.G.E. Consortium/LLNL at:  http://image.llnl.gov  plate: LLAM14017 row: n column: 21  High quality sequence stop: 543.  Location/Qualifiers  1 040	Contact: Robert Strausberg, Ph.D.		National Institutes of Health, Mammalian Gene Collection (MGC)	NIH-MGC http://mgc.nci.nih.gov/.	1 (bases 1 to 940) .	Rodentia; Sciurognathi; Muridae	Eukaryota; Metazoa;	M Mus musculus	house mouse.	EST.	BQ937887.1 GI:22353365		IMAGE:6476852 5', mRNA sequence.	AGENCOURT_8951807	BQ937887 940 bp mRNA linear EST 21-AUG-2002	

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CDNA clone set

Unpublished (2002)

Contact: Yong Qian

Laboratory of Genetics

National Institute on Aging/National Institutes of Health

333 Cassell Drive, Suite 3000, Baltimore, MD 21224-6820, USA

Email: cdna@lgsun.grc.nia.nih.gov

This clone set has been freely distributed to the community.

visit http://lgsun.grc.nia.nih.gov/cDNA/NIA_7_4k.html for de

Plate: H4067 row: A column: 01

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                                                                                                                                                                                                                                                                                                                                                                                                                                  VanBuren, V., Piao, Y., Dudekula, D.B., Qi, P.R., Stagg, C.A., Bassey, U., Aiba, K., Luo, A.G. and Ko, M.S.H.
Assembly, verification, and initial ann CDNA clone set
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
Eukaryota; M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            house mouse
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Similarity 100.0%;
0; Conservative
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                87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:10090"
/clone="IMAGE:6476852"
/clone_lib="NCI_CGAP_Co24"
/clone_lib="NCI_CGAP_Co24"
/lab_host="DH10B (Tl phage-resistant)"
/note="Organ: colon; Vector: pcMV-SPORT6; Site_1: No Site_2: Sall; Cloned unidirectionally. Primer: Olignerage insert size 1.6 kb. Constructed by Life Technologies. Note: this is a Note of Site_1 CGAP Library."
9 a 277 c 288 g 200 t 6 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ø
               þ
             clone is
than 20
a 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eutheria; 1 to 537)
           /note-"Vector: pSPORT1; Site_1: clone is among a rearrayed set than 20 cDNA libraries."

162 c 166 g 122 t
                                                                                                                    /strain="C57BL/6"
/db_xref="niaEST:H4067A01-5
/db_xref="taxon:10090"
/clone="H4067A01"
/clone_lib="NIA Mouse 7.4K
                                                                         /dev_stage="mixed"
/lab_host="DH108"
                                                                                                                                                                                                  organism-"Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Mus musculus/strain="FVB/N"
                                                                                                          'sex="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NIA Mouse 7.4K cDNA Clone, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GI:21461753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chordata;
Rodentia;
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Pred.
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Mismatches 0;
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Sciurognathi; Muridae;
                                                                                                                      7.4K CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set Mus musculus cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Qian, Y., Carter, M.G., Martin
., Hamatani, T., Kargul, G.J.,
                                            : SalI;
                                                                                                                        Clone
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                                         I; Site_2:
,407 clon-
                                                                                                                        Set
                                           clones
                                                                                                                                                                                                                                                                                                          community. Please tml for details.
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Murinae; Mus
                                                            NotI;
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BB849097
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                                                                                                                                                       Email: genome-res@gsc.riken.go.jp,

URL:http://genome.gsc.riken.go.jp/

Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.

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Please visit our web site (http://genome.gsc.riken.go.jp) for further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The Institute 1-7-22 Suehiro-cho, Tel: 81-45-503-9222
Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Akimura, T., Arakawa, T., Carninci, P., Furuno, M., Hanagaki, T., Hayatsu, N., Hiramoto, K., Hiraoka, T., Hirozane, T., Imotani, K., Ishii, Y., Ito, M., Kawai, J., Kojima, Y., Konno, H., Kouda, M., Matsuyama, T., Nakamura, M., Nishi, K., Nomura, K., Numasaki, R., Okazaki, Y., Okido, T., Saito, R., Sakai, C., Sakai, K., Sakazume, N., Sasaki, D., Sato, K., Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Suzuki, H., Tagawa, A., Takahashi, F., Takaku-Akahira, S., Tanaka, T., Tomaru, A., Toya, T., Watahiki, A., Yasunishi, A., Muramatsu, M. and Hayashizaki, Y. RIKEN Encyclopedia of Mouse Full-length cDNAs (Akimura, T., et al. 2001)
Unpublished (2001)
Contact: Yoshihide Hayashizaki
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1 (bases
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BB849097 RIKEN full-length enriched, adult cDNA clone F930006G02 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="F930006G02"
/clone_lib="RIKEN full
                                                                                                                                 Location/Qualifiers
                                                                                    ∕organism="Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 to 424)
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 adult
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Sathe,G.,
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1 (bases
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                                                                                                                                                                         32;
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Identification and initial characterization sequenced tags (ESTs) each from adult human osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001) 21482651
Contact: Sanjay Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   709 Swedeland Road
Tel: 610-270-7245
Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GlaxoSmithKline
709 Swedeland R
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Eukaryota; M
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9; Conservative
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llarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sanjay_kumar-1@gsk.com
imer: T7.
                                                                                                                                                                                                                                                                          /tissue_type="cartilage"
/lab_host="E.coli DH10 B'
/note="Vector: pSPORT I;
Directional"
a 139 c 118 g 65
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sex=male), (tissue_type=whole body, dev_stage=9 days
embryo, sex=mixed), (tissue_type=lung, dev_stage=13 da
embryo, sex=mixed)
124 c 132 g 103 t
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/dev_stage="adult"
/note="pooled tissues ;
dev_stage=16 days neona
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/db_xref="taxon:9606"
/clone_lib="HOA (Human O
/tione_name of the content of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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Mui, P., Agarwal, P.,
                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Homo
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type=cerebellum, dev_stage=0 day neonate
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Primates;
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al,P., Badger,A.M., Le
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Lee,J.C.,
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Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries

Costeoarthr. Cartil. 9 (7), 641-653 (2001)

E 21482651

Contact: Sanjay Kumar
UW2109

GlaxoSmithKline
709 Swedeland Road, P.O. Box 1539, King of Prussia, PA 19406, US/Tel: 610-270-7245
Fax: 610-270-5598

Email: sanjay_kumar-1@gsk.com
Seq primer: T7.
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BG538695

BG538695.1 GI:13530928

EST.

human.

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutemannalia; Eutheria; Primates; Catarrhini; Hominidae; Homannalia; Eutheria; Primates; Catarrhini; Hominidae; Homannalia; Eutheria; Primates; Catarrhini; Hominidae; Homannalia; Choses 1 to 618)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
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found http:/
                                                                                                                                                                                                                                                                                                                    BG538695 618 bp r
602566932F1 NIH_MGC_77 Homo sapiens
         Email: cgapbs-r@mail.nih.gov
Fissue Procurement: CLONTECH Laboratories, Inc.
cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
ctp://image.llnl.gov
                                                                                                                                                  IH-MGC http://mgc.nci.nih.gov/.
itional Institutes of Health, Mammalian Gene Collection (MGC)
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/db_xref="taxon:9606"
/clone_lib="HOA (Human Osteoarthritic
/tissue_type="cartilage"
/lab_host="E.coli DH10 B"
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IMAGE:4691574 5',
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Sathe,G.,
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High qu
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Identification and initial characterization sequenced tags (ESTs) each from adult human osteoarthritic cartilage cDNA libraries
Osteoarthr. Cartil. 9 (7), 641-653 (2001) 21482651
Contact: Sanjay Kumar
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32; Conservative
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Fax: 610-270-5598
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709 Swedeland R
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/db_xref="taxon:9606"
/clone="IMAGE:4691574"
/clone="IMAGE:4691574"
/clone="IMAGE:4691574"
/clone="Ib="NIH_MGC_77"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
SfiI (ggccgctcggcc); Site_2: SfiI (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
5'-ATTCTAGAGGCCGAGGCGGCCGACATG-GT(30)BN-3' (where B = A,
C, or G and N = A, C, G, or T) Average insert size 1.9
kb (range 0.5-4.0 kb). 12/15 collonies contained inserts
by PCR. This library was enriched for full-length clones
and was constructed by Clontech Laboratories (Palo Alto,
CA). Note: this is a NIH_MGC Library."
                                                                                                                                                                         sanjay_kumar-l@gsk.com
lmer: T7.
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/clone_lib="HNC (Human No
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                                                      /tissue_type="cartilage"
/lab_host="E.coli DH10 B
/note="Vector: pSPORT I;
Directional"
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Mui, P., Agar
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Agarwal,P., Bad
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Badger, A.M., Le
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1869 row: g column: 24
                                                                                                                                                                                                                                                                                                                        375
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AL555144
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Mammalia;
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/clone=lib="NIH_MGC_42"
/tissue_type="epithelioid carcinoma cell line"
/tissue_type="epithelio"
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Eutheria; Primates;
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Eukaryota;
Mammalia; I
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Email:
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                                                                                                                                                                                                                                                                                                                                                                    human.
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National Institutes of Health, Mammalian
Unpublished (1999)
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Mammalia; Eutheria;
1 (bases 1 to 750)
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                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D Email: cgapbs-r@mail.nih.gov Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                            1 (bases 1 to NIH-MGC http://
                                                                                                                                                                                                                                                                                                                                                                                      EST.
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larity 100.0%;
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/db_xref="taxon:9606"
/clone="CS0DK007Y021"
/clone_lib="LTI_NFL006_PL2"
/tissue_type="placenta"
/db_xref="taxon:9606"
/clone="IMAGE:5446794"
/clone_lib="NIH_MGC_AO"
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Primates;
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Catarrhini;
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IMAGE:5446794 5',
                                                                                                                                                                                                                                                                                                                           Euteleostomi; Homo.
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B1826781.1 GI:15938331

EST.

human.

Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 800)

S NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

L Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.
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Plate: LLAM11420 row: g column: 18
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603077268F1 NIH_MGC_119 F
mRNA sequence.
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  Similarity 100.0%; 2; Conservative
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Llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                            /db_xref="taxon:9606"
/clone="IMAGE:5169065"
/clone_lib="NIH_MGC_119"
/tissue_type="medulla"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
                                                                                                                 site_2: EcoRV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics +racking this is a war.
                                       5.0%;
                                                                                           a NIH_MGC Library.
c 264 g 136
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RESULT 12
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                                                BI825652
603072631F1 NIH_MGC_119 H
mRNA sequence.
BI825652
BI825652.1 GI:15937202
EST.
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603039845F1 NI
mRNA sequence.
BI822142
BI822142.1 GI
EST.
  Eukaryota;
Mammalia; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Straus.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution informat
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Mammalla; E
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Similarity 100.0%;
32; Conservative
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National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                       Homo sapiens
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                                                                                                                                                                                                                                                                                                 134
                                                                                                                                                                                                                                                                      source anonymous pool of 6 male brains, age range 23-27; male lung, age 27; and 1 male testis, age 69. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.8 kb, insert size range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C Gruber (Invitrogen). Research Genetics tracking code 021. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                            / Janism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5180650"
/clone_lib="NIH_MGC_115"
/lah host="rorr"
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  Metazoa;
Eutheria;
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Primates;
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Craniata; Vertebrata; Catarrhini; Hominidae;
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Eukaryota; Metazoa; Chordata; Crani
Mammalia; Eutheria; Primates; Catar
1 (bases 1 to 916)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mamm
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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BI457367
BI457367.1 GI
EST.
human.
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11408 row: n column: 07
High quality sequence stop: 877.
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mal
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1870 row: a column: 23 High quality sequence stop: 778.
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603185689F1 NIH_MGC_42 Homo sapiens
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Similarity 100.0%;
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Location/Qualifiers

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              ity sequence stop: 778. Location/Qualifiers
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/clone_lib="NIH_MGC_119"
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lab_host="DH108"
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Pred. No. 0.00017;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov

Tissue Procurement: ATCC,
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1833 row: m column: 18
High quality sequence start: 28
High quality sequence stop: 756.
Location/Qualifiers
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Mammalia; Eut
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg,
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                          /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5019065"
/clone=lib="NIH_MGC_42"
/tissue_type="epithelioid carcinoma cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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/tissue_type="epithelioid carcinoma cell line"
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SUMMARIES

## Result No. Scor 218 218 166 117 000000011111112222277222222222222Query Match æ 137964 137964 1741 1734 1734 61072 216757 1739 1749 Leng AC126895 AC095418 AC095418 AC126895 AF210324 AR210325 AF100778 AF1100778 AF126063 AF210337 AR210337 AR210338 AX076919 AX464186 AF083500 AR210322 AR210323 AF074604 AF100780 BC017782 AL139352 AL139352 AC112914 AC112856 AC1122102 AL646068 AC1122102 AL671882 AL646068 AC1122102 AL671882 AL671882 AL671882 AL6719257 AB020659 AB020659 AB020659 AB0205541 AC195541 ALIGNMENTS AC126895 Rattus no AC095418 Rattus no AC126895 Rattus no AC126895 Rattus no AC126895 Rattus no AC12689581 Rattus no AF259981 Rattus no AF259981 Rattus no AR210324 Sequence AF100778 Mus muscu AF126063 Mus muscu AF126063 Mus muscu AF126063 Mus muscu AR210337 Sequence AX076919 Sequence AX076919 Sequence AF083500 Homo sapi AR210322 Sequence AF074604 Homo sapi AC110782 Homo sapi BC017782 Homo sapi AC110904 Mus muscu AC116351 Homo sapi AC119914 Mus muscu AC112856 Rattus no AL646068 Ralstonia AC119914 Mus muscu AC1123102 Rattus no AL671882 Mouse DNA AL646068 Ralstonia AC1133637 Homo sapi AB033535 Oryza sat BC019257 Homo sapi Description

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pieces.	H23(	137964 bp DNA	

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REFERENCE AUTHORS	CRGANLSM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	AC126895	RESULT 1
Rattus.  1 (bases 1 to 137964)  Muzny,D.M., Adams,C., Adio-Oduola,B., Ali-osman,F.R., Allen,C.,	Kattus norvegicus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;	Rattus norvegicus.	HTG; HTGS_PHASE1.	AC126895.1 GI:21724040	AC126895	***, 49 unordered pieces.	H230-301E4, *** SI	AC126895 137964 bp DNA linear HTG 24-JUL-200		

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Alsbrooks, S. L., Amaratunge, H.C., Are, J. R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Boule, S., Brievam, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burrell, K. L., Byrd, N.C., Carter, M., Cavazos, S. R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Ghristopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Ghristopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Ghristopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Baderich, D.A., David, R., David, R
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3 (bases 1 to 137964)
Worley, K.C.
Direct Submission
Submitted (24-JUL-2002) F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (24-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA

Center: Baylor College of Medicine Center code: BCM
Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu/
                                                (see
                                                                                                                                                           NOTE:
                                                                                                                                                                                                                                            Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: Assembly program: Phrap; version Consensus quality: 89474 bases at Consensus quality: 93422 bases at Consensus quality: 936506 bases at
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Center project name: GZHG
Center clone name: CH230-301E4
------ Summary Statistics
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Stimated insert size may differ from sequence length, http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_ehts is a 'working draft' sequence. It currently of 49 contigs. The true order of the pieces
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Genetics, Baylor College
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank\_draft\_data.html).

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Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 185950 bases at least Q40 Consensus quality: 190362 bases at least Q30 Consensus quality: 193076 bases at least Q20
                                                                                                                                                                                                                                   Center ....
Web site: http://www....
Contact: hgsc-help@bcm.tmc.edu
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REFERENCE AUTHORS

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Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Davila, M.L., Davis, C., Davy-Carroll, L., Ding, Y., Dinh, H.H., Delanney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delanney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delanney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Gill, R., Garler, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Garbis, A., Gao, J., Garcia, A., Garner, T., Garza, M., Gill, R., Gorrell, J.H., Guevara, W., Gunarratne, P., Hale, S., Hamilton, K., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Hemrandez, J., Homsi, F., Howard, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Karlsson, E., Hober, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Liu, W., Loulseged, H., Lozado, R.J., Lu, M., Martindale, A., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Maheshwari, M., Morris, S., Moser, M., Newtson, J., Newtson, M., Martindale, A., Martinez, E., Put, L.L., Quiles, M., Ren, Y., Peters, L., Peters, L., Pace, A., Payton, B., Peery, J., Perez, L., Peters, L., Pace, A., Payton, B., Peery, J., Perez, L., Peters, L., Put, L.L., Quiles, M., Ren, Y., Sodergren, E., Sonaike, T., Sparks, A., Kolles, M., Thomas, N., Thomas, S., Wulliamson, A., Warren, R., Washington, C., Watlington, S., Worley, K., Wang, O., Watlington, S., Worley, K., Warley, K., Warley, K., Wang, O., W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (10-JUL-2002) Human Genome Sequencing Ceof Molecular and Human Genetics, Baylor College of Baylor Plaza, Houston, TX 77030, USA

3 (bases 1 to 137964)

Worley, K.C.

Direct Submission
Submitted (24-JUL-2002) Human Genome Sequencing Ceof Molecular and Human Genetics, Baylor College of Baylor Plaza, Houston, TX 77030, USA

Center: Baylor College of Medicine
Center code: BCM

Center code: BCM
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Direct Submission
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Contact: hgsc-help@bcm.tmc.edu
Center project Information
Center project name: GZHG
Center clone name: CH230-301E4
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COMMENT

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TITLE JOURNAL

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Consensus quality: 96506 bases at least Q20
NOTE: Estimated insert size may differ from sequence length
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank\_draft\_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 49 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.

1 1056: contig of 1056 bp in length
1157 1156: gap of unknown length

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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

E 1 (bases 1 to 1741)
S Zhang, R., Averboukh, L., Zhu, W., Zhang, H., Jo, H., Dempsey, P.J.,

Coffey, R.J., Pardee, A.B. and Liang, P.

Identification of rCop-1, a new member of the CCN protein family;

as a negative regulator for cell transformation

L Mol. Cell. Biol. 18 (10), 6131-6141 (1998)

9 942130
E 2 (bases 1 to 1741)
S Liang, P.

Direct Submission
L Submitted (24-APR-2000) Cell Biology, Vanderbilt-Ingram Cancer

Center, 649 MRB II, Nashville, TN 37232, USA
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                                               Mus musculus.

Mus musculus

Eukaryota; Metazoa; Cl

Mammalia; Eutheria; Rc

1 (bases 1 to 1734)

Pennica, D., Swanson, T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
Unclassified.
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Sequence
AR210325
AR210325.
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Mus muscu
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Botstein, D.A., Cohen, R.L., Goddard, A.D., Gurney, A
Lawrence, D.A., Levine, A.J., Pennica, D., Roy, M.Ann
WISP polypeptides and nucleic acids encoding same
Patent: US 6387657-A 18 14-MAY-2002;
                   Pennica, D., Swanson, T.A., Welsh, J.W., Roy, M.A., Lee, J., Brush, J., Taneyhill, L.A., Deuel, B., Lew Cohen, R.L., Melhem, M.F., Finley, G.G., Quirke, P. Hillan, K.J., Gurney, A.L., Botstein, D. and Levin,
                                                                                                                                                                                                                                                                                                                    90;
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                                                                                                                               Mus musculus connective (Wisp2) mRNA, complete of AF100778
AF100778.1 GI:4028578
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          Gurney, A.L., ce members of
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nley,G.G., Quirke,P., Goddard,A.D., otstein,D. and Levine,A.J.
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Location/Qualifiers
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/gene="Wisp2"
/codon_start=1
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/translation="MRGNPLIHLLAISFLCILSMVYSQLCPAPCACPWTPPQCPPGVP
/translation="MRGNPLIHLLAISFLCILSMVYSQLCPAPCACPWTPPQCPPGVP
/translation="MRGNPLIHLLAISFLCILSMVYSQLCPAPCACPWTPPQCPPGVP
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NGRRYLDGCTFKPNCRVLCRCDDGGFTCLPLCSEDVRLPSWDCPRPRRIQVPGRCCPE
WVCDQAVMQPAIQPSSAQGHQLSALVTPASADGPCPNWSTAWGPCSTTCGLGIATRVS
NQNRFCQLEIQRRLCLSRPCLASRSHGSWNSAF"
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/tissue_type="mammary"
/note="transformed by Wnt-1"
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Pred. No.
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~1one RP23-161B3 on
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95 (25), 14717-14722
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For further details see http://www.chori.org/bacpac/home.htm
410
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Mus musculus chromosome 2 clone RP2
PROGRESS ***, 21 unordered pieces.
AL669906

AL669906.5 GI:21955520

HTG; HTGS_PHASE1; HTGS_CANCELLED.
Mus musculus.
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Eukaryota; Me
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/db_xref="taxon:10090"
/chromosome="2"
/clone="RP23-161B3"
/clone_lib="RPCI-23"
a 15368 c 15033 g 15260
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Eutheria; Rodentia;
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Pred. No.
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      /note="assembly_fragment_chain:139772...46360
                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
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/clone_lib="RPCI-23"
                                                                                                                                                                                                                                                      vector_
                                                                                                                                                                 4101. .15267
/note="assembly_fragment:00814
fragment_chain:1"
15368. .34970
                                                                                           fragment_c
35071. .39
                                                                                                                                                                                                                                                                                                              clone_end:SP
                                                                                     /note="assembly_fragment:00812
fragment_chain:1"
35071. .39671
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clone_end:SP6
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___...ω1y_fragment:03544
t_chain:1"
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RESULT 10
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AF126063
Mus musculus o
(Ctgfl) mRNA,
AF126063
                                                                                                                                                                                                            90;
                                                                                                                                                                                                           n 10.2%;
Similarity 100.0%;
90; Conservative
                                                                                                      GGGCCTGGTTTGTCAGCCTGGGGCAGGCCC
                                                                                                                                                      AGTGTGTGCACGGAGGCTGGGGGAGTCCTGCGACCACCTGCATGTCTGCGACCCCAGCCA
                                                                                                                                                                                                                                                                             6425
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/note="assembly_fragment:02352
fragment_chain:5"
151673. .169686
/note="assembly_fragment:02207
fragment_chain:5"
169787. .200095
/note="assembly_fragment:02849
fragment_chain:6"
200196. .213877
/note="assembly_fragment:01052
fragment_chain:6"
213978. .216757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="assembly_fragment:01170 fragment_chain:3" 98702. .105522 /note="assembly_fragment:02829 fragment_chain:3" 105623. .119773 /note="assembly_fragment:01476 fragment_chain:4" 119874. .134552 /note="assembly_fragment:00505 fragment_chain:4" 134653. .141202 /note="assembly_fragment:01951 fragment_chain:4" 141303. .145808 /note="assembly_fragment:02637 fragment_chain:5" 145909. .151572
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51573 c 51904 g
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fragment_chain:2"
84736. .87399
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fragment_chain:1"
46461: .49809
/note="assembly_fragment:03410
fragment_chain:2"
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clone_end:T7
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fragment_chain:3"
87500. .98601
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fragment_chain:2"
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fragment_chain:2"
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           complete
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                                                                                                                                                                                                          0;
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Pred. No.
             1739 bp
e tissue ç
cds.
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                         growth
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5.6e-37;
0;
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                         mRNA linear
th factor-like
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                      ROD 12-OCT-1999 protein precursor
                                                                                                                                                                                                            0;
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                                                                                                                                                                                  469
                                                                                                                                                         166509
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KEYWORDS
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RESULT 11
AR210337
LOCUS
DEFINITION
ACCESSION
VERSION
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KEYWORDS
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Best Local S
Matches 80
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AR210337.
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umar, S., Hand, A.T., Connor, J.R., Dodds, R.A., Ryan, P.J.,
rill, J.J., Fisher, S.M., Nuttall, M.E., Lipshutz, D.B., Zou, C.
wang, S.M., Votta, B.J., James, I.E., Rieman, D.J., Gowen, M. ar
ee, J.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          irect Submission
ubmitted (04-FEB-1999) Bone & Cartilage Biology, UW 210
mitherline Beecham, 709 Swedeland Rd., King of Prussia,
                                                                                                                                                                                                                             Similarity
0; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9287915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dentification and cloning of a connective tissue gactor-like cDNA from human osteoblasts encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             us musculus.
                                            R210337
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                                                                                                                                                                                                                         9.1%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="lung"
1. .1739
/gene="Ctgfl"
242. .997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chem.
                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="similar to the Mus musculus WISP-2 protein encoded by the sequence presented in GenBank Accession Number AF100778; putative growth factor; CTGF-L; contains IGF binding (IGFBD), Von Willebrand Factor type C (VWC) repeat and thrombospondin type I (TSP1) domains; member of the CCN (CTGF/Cyr61/Nov) family; lacks the fourth carboxy-terminal (CT) domain present in other members of the CCN family"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                s 1 to 1739)
and Zou,C.
                                                                                                                                                                                                                                                                                            /protein_id="AAD18058.1"
/db_xref="GI:4337060"
/translation="MRGNPLIHLLAISFLCILSMVYSQLCPAPCACPWTPPQCPPGVP
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LVLDGCGCCRVCARRLGESCDHLHVCDPSQGLVCQPGAGPSGRGAVCLFEEDDGSCEV
NGRRYLDGETFKPNCRVLCRCDDGGFTCLPLCSEDVRLPSWDCPRPRRIQVPGRCCPE
WVCDQAVMQPAIQPSSAQGHQLSALVTPASADGPCPNWSTAWGPCSTTCGLGIATRVS
NQNRFCQLEIQRRLCLSRPCLASRSHGSWNSAF"
480 c 489 g 395 t
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                                                                                                                                                                                                                                                                                                                                                                                                                         /product="connective tissue growth factor-like
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   GI:21512542
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Rodentia;
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Pred. No.
                              738 bp
US 6387657
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                                                                                                                                                                                                                        DB 10; 4
3.2e-31;
0;
                                                                                                                                                                                                                                                        Length 1739;
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                                            20-JUN-2002
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AR210338
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1 (bases 1 to 738)
Botstein,D.A., Cohen,R.L., Goddard,A.D., Gurney
Lawrence,D.A., Levine,A.J., Pennica,D., Roy,M.A
WISP polypeptides and nucleic acids encoding sa
Patent: US 6387657-A 38 14-MAY-2002;
Location/Qualifiers
1. .738
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Sequence
AR210338
AR210338.
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Sequence
AX076919
AX076919.
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                                                                  Eukaryota; Metazoa; Choromammalia; Eutheria; Prim; 1 (bases 1 to 1266)
Botstein, D., Goddard, A.,
                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 841)
Botstein, D.A., Cohen, R.L., Goddard, A.D., Gurney, A
Lawrence, D.A., Levine, A.J., Pennica, D., Roy, M.Ann
WISP polypeptides and nucleic acids encoding same
Patent: US 6387657-A 39 14-MAY-2002;
       Polypeptidic compositions and methods Patent: WO 0105836-A 31 25-JAN-2001; Genentech, Inc. (US)
Location/Qualifiers
1. .1266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
32; Conser
                                                           Botstein,D.,
Wood,W.I.
                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
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                                                                                                            Homo sapiens
                                                                                                                       human
                                                                                                                                                                                                                                                                                                                                                                                                            Unclassified.
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                                                                                                                                                                                                                                                                3.6%;
Similarity 100.0%;
32; Conservative
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larity 100.0%;
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1. .841
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272 c 238 g
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organism="Homo
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d Wood,W.I
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2 (bases
Kumar,S.
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SmithKline
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                                                    Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1283)

Kumar,S., Hand,A.T., Connor,J.C., Dodds,R.A., Ryan,P.J.,

Trill,J.G., Fisher,S.M., Slemmon,J.R., Lipshutz,D.B.,

Bartholomew,V., James,I.E., Rieman,D.J., Gowen,M. and Lee,J.C.

Identification and cloning of CTGF-L from human osteoblasts, a novel cysteine rich protein containing an IGF binding domain Bone 23 (5), S240 (1998)

2 (bases 1 to 1283)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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Baker,K.P., Beresini,M., Deforge,L., Desnoyers,L., Filvaroff,E.,
Gao,W.Q., Gerritsen,M.E., Goddard,A., Godowski,P.J., Gurney,A.L.,
Sherwood,S., Smith,V., Stewart,T.A., Tumas,D., Watanabe,C.K.,
Wood,W.L. and Zhang,Z.
Secreted and transmembrane polypeptides and nucleic acids encoding same
Patent: WO 0140466-A 319 07-JUN-2001;
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db_xref="taxon:9606"
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Beecham, 709
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/db_xref="taxon:9606"
/chromosome="20"
/map="20q12-q13"
/cell_type="primary osteoblast"
9. .761
                                                                                                   Location/Qualifiers
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## ALIGNMENTS

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acid sequences encoding rat heparin-induced CCN-like protein, methods to identify modulators or in diagnostic applications
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induced CCN-like protein; HICP; cell-associated activity; ss; scular disorder; aberrant cell proliferation; fibrotic disorder

Rat HICP

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Agents that stimulate or inhibit HICP protein activity or sion, antisense HICP nucleic acid molecules and HICP antibodies, used to modulate cell-associated activity. HICP modulators can be treat disorders characterized by aberrant HICP protein activity ession. Probes capable of hybridizing to HICP mRNA or antibodies of for HICP can be used to detect HICP activity in a biological HICP can be used to treat disorders, such as a cardiovascular or c disorder, characterized by aberrant cell proliferation. The sequence represents the coding sequence of rat HICP.
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The invention provides a rat heparin-induced CCN-like protein (HICP) protein. Agents that stimulate or inhibit HICP protein activity or expression, antisense HICP nucleic acid molecules and HICP antibodies, can be used to modulate cell-associated activity. HICP modulators can used to treat disorders characterized by aberrant HICP protein activit or expression. Probes capable of hybridizing to HICP mRNA or antibodic specific for HICP can be used to detect HICP activity in a biological
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TGCTCAACCACCTGTGGGCTGGGCAT
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induced CCN-like protein; HICP;
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P-PSDB;
                                                                                             Heparin-induced CCN-like cardiovascular disorder; Von Willebrand C domain;
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RESULT 6
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testicular
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tissue-gro
kidney dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention provides a rat heparin-induced CCN-like protein (HICP) protein. Agents that stimulate or inhibit HICP protein activity or expression, antisense HICP nucleic acid molecules and HICP antibodies, can be used to modulate cell-associated activity. HICP modulators can be used to treat disorders characterized by aberrant HICP protein activity or expression. Probes capable of hybridizing to HICP mRNA or antibodies specific for HICP can be used to detect HICP activity in a biological sample. HICP can be used to treat disorders, such as a cardiovascular or fibrotic disorder, characterized by aberrant cell proliferation. The present sequence represents a cDNA encoding the Von Willebrand C (VWC) a domain of the HICP polypeptide.
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97;
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                                                                                                                                                                         sorder; bone-related disorder; osteoporosis; trauma; e tissue disorder; catabolic state; inflammation; r-related disorder; angiogenesis; immunological disor
                                                                                                                                                                                                                   uced secreted protein; WISP-1; WISP-2; WISP-3; CTGF; tumour e tissue growth factor; cancer; melanoma; arteriosclerosis; lymphoid malignancy; haematopoiesis-related disorder; owth disorder; skin disorder; desmoplasia; fibrotic lesion;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences encoding rat heparin-induced CCN-like ods to identify modulators or in diagnostic appliance.
                                                                                                                                                                                                                                                                                            protein complementary nucleotide sequence SEQ ID NO:18.
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97US-0063704.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Products from the present invention can be used to treat WISP-related disorders such as breast, ovarian, and colon cancer or melanoma. The products can be used to treat arteriosclerosis. The products can also used to treat other diseases e.g. benign and malignant tumours, leukaemia and lymphoid malignancies, neuronal, glial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, stromal, and chisorders, skin disorders, haematopoiesis-related disorders, tissue-growt disorders, skin disorders, desmoplasia, fibrotic lesions, kidney disorders, bone-related disorders such as osteoporosis, trauma such as until control of the states, testicular-related disorders, and inflammatory, angiogenic and immunologic disorders including arteriosclerosis. The products can also be used for detection and diagnosis especially of individuals with neoplastic cell growth or proliferation. The products can be used in the production of transgenic or knock-out animals.

C Antibodies can be used to induce death in WISP-1, 2 or 3 overexpressin
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Best Local S
Matches 90
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Antibodies
cells.
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                                                                                                                                                                                 WNT-1 induced secreted protein; WISP-1; WISP-2; WISP-3; CTGF; connective tissue growth factor; cancer; melanoma; arterioscle leukaemia; lymphoid malignancy; haematopoiesis-related disordetissue-growth disorder; skin disorder; desmoplasia; fibrotic kidney disorder; bone-related disorder; osteoporosis; trauma; connective tissue disorder; catabolic state; inflammation; testicular-related disorder; angiogenesis; immunological disorder
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90; Conservative
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haematopoiesis-related disorder;
sorder; desmoplasia: fire...
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29-OCT-1997;
03-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                  Leukaemia and lymphoid malignancies, neuronal, glial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, stromal, and blastocoelic disorders, haematopoiesis-related disorders, tissue-growth disorders, skin disorders, desmoplasia, fibrotic lesions, kidney disorders, bone-related disorders such as osteoporosis, trauma such as burns, incisions, and other wounds, connective tissue disorders, catabolic states, testicular-related disorders, and inflammatory, angiogenic and immunologic disorders including arteriosclerosis. The products can also be used for detection and diagnosis especially of individuals with neoplastic cell growth or proliferation. The products can be used to induce death in WISP-1, 2 or 3 overexpressing
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P-PSDB;
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AAY17651.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and 3. The novel WISP polypeptides, designated WISP-1, WISP-2 and 3. The novel WISP polypeptides, designated WISP-1, WISP-2-3 have homology to connective tissue growth factor (CTGF). from the present invention can be used to treat WISP-related such as breast, ovarian, and colon cancer or melanoma. The can be used to treat arteriosclerosis. The products can also be
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97US-0063704.
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                                                                                      HICP; cell-associated activity; ss;
cell proliferation; fibrotic disorder;
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04-OCT-2000;
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P-PSDB;
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nt invention relates to single exon nucleic acid probes for human gene expression in a sample derived from human heart. equence is one such probe. The probes may be used for g, measuring and displaying gene expression in samples derive human heart via microarrays. By measuring gene expression, the useful for predicting, diagnosing, grading, staging,
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                                                                                                                                                                 OLECULAR DYNAMICS
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                                                                                         The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the invention.
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Note: The sequence data for this patent did specification, but was obtained in electronicat figure.int/pub/published_pct_sequences.
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Note: The sequence data for this patent did not of the printed specification, but was obtained format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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TITLE OF INVENTION: WISP POLYPEPTIDES AN FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,14
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,7(
EARLIER APPLICATION NUMBER: US 60/073,61
EARLIER APPLICATION NUMBER: US 60/073,61
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: US 60/081,65
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
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TITLE OF INVENTION: WISP POLYPEPT
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60
EARLIER FILING DATE: 1997-10-29
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EARLIER APPLICATION NUMBER: US 60
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JRRENT APPLICATION NUMBER: US/09/182,145B

JRRENT FILING DATE: 1998-10-29

ARLIER APPLICATION NUMBER: US 60/063,704

ARLIER APPLICATION NUMBER: US 60/073,612

ARLIER FILING DATE: 1998-02-04

ARLIER APPLICATION NUMBER: US 60/081,695

ARLIER APPLICATION NUMBER: US 60/081,695

ARLIER FILING DATE: 1998-04-14

UMBER OF SEQ ID NOS: 156
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ICANT: Roy, Margaret Ann
ICANT: Wood, William I.
E OF INVENTION: WISP POLYPEPTIDES AND PREFERENCE: P1176R2
ENT APPLICATION NUMBER: US/09/182,145B
ENT FILING DATE: 1998-10-29
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Pennica, Diane
Roy, Margaret Ann
Wood, William I.
NVENTION: WISP POLYPEPTIDES A
RENCE: P1176R2
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Gurney, Austin L.
Hillan, Kenneth J.
Lawrence, David A.
Levine, Arnold J.
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US-09-182-145-14/c
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                                                                                                        GENERAL
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EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/CURRENT FILING DATE: 1998-10-29 EARLIER APPLICATION NUMBER: US EARLIER FILING DATE: 1997-10-29 EARLIER APPLICATION NUMBER: US
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APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES
FILE REFERENCE: P1176R2
                                                                                  APPLICANT:
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                                 Cohen, Ro
Goddard,
            Gurney,
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Hillan, Kenneth J.
Lawrence, David A.
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                                                                                David A
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Pred. No.
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APPLICANT: ENDEGE, WILSON O., ET AL.

TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION

TITLE OF INVENTION: PRODUCTS: II

FILE REFERENCE: CCDNA-260XX

CURRENT APPLICATION NUMBER: US/09/385,982

CURRENT FILING DATE: 1999-08-30

EARLIER APPLICATION NUMBER: 09/328,111

EARLIER APPLICATION NUMBER: 60/117,393

EARLIER APPLICATION NUMBER: 60/117,393

EARLIER APPLICATION NUMBER: 60/098,639

EARLIER APPLICATION NUMBER: 60/098,639

EARLIER FILING DATE: 1998-08-31

NUMBER OF SEQ ID NOS: 544
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Db
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US-09-385-98
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LENGTH: 51
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PPLICANT: Wood, William I.
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PPLICANT: Wood, William I.
PPLICATION NUMBER: US/09/182,145B
PRAILER APPLICATION NUMBER: US 60/063,704
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER: OF SEQ ID NOS: 156
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NAME/KEY:
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TYPE: DNA
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R INFORMATION: n = 15-982-220
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Similarity 100.0%;
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US-99-149-476-225/c
Sequence 225, Application US/09149476
Sequence 225, Application US/091494
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: 60/040,16:
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,601
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ILING DATE: 1997-05-23
PPLICATION NUMBER: 60/043,670
ILING DATE: 1997-04-11
PPLICATION NUMBER: 60/056,632
ILING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,632
ILING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,876
ILING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,876
ILING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,875
ILING DATE: 1997-08-22
ILING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,875
ILING DATE: 1997-08-22
                                                    : Brenda F. Baker
: Lex M. Cowsert
!NVENTION: ANTISENSE MODULATION OF S
RENCE: RTS-0026
PPLICATION NUMBER: US/09/213,768
!LING DATE: 1998-12-17
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LING DATE: 195,
PLICATION NUMBER: 60/0-23
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LICATION NUMBER: 60/056,908
ING DATE: 1997-08-22
LICATION NUMBER: 60/048,964
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ING DATE: 1997-05-23
LICATION NUMBER: 60/047,590
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985664
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LICATION NUMBER: 60/1
ING DATE: 1997-09-05
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ilarity 100.0%;
Conservative 0
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Pred. No.
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6.2;
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Sequence 2, Application US/09213768; Patent No. 5985664; GENERAL INFORMATION:
APPLICANT: Brenda F. Baker; APPLICANT: Lex M. Cowsert; TITLE OF INVENTION: ANTISENSE MODULATION OF FILE REFERENCE: RTS-0026; CURRENT APPLICATION NUMBER: US/09/213,768; CURRENT FILING DATE: 1998-12-17; NUMBER OF SEQ ID NOS: 47
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TITLE OF INVENTION: No. 6436703el Nucleic
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 790CIP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
                                                                                                                                                                                                            Query Match
Best Local S
Matches 19
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Matches
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LOCATION: (136
-09-213-768-1
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                                                                                                                 09-213-768-2/c
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                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID
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                                                                                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: (130
09-668-680-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: FEATURE:
                                                                                                                                                                                                                                                                                         ORGANISM: Homo FEATURE:
                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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| Similarity 100.0%;
| 19; Conservative
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Similarity 100.0%;
19; Conservative
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Xue, Aidong J
Xu, Chongjun
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Liu, Chenghua
Zhou, Ping
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Zhang, Jie
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                                               MODULATION OF
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Pred. No. 6.2
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NAME: Daniel W. Collins
REGISTRATION NUMBER: 31,912
REFERENCE/DOCKET NUMBER: 5381.US.P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 937-6365
TELEFAX: (708) 938-2623
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 2949 base pairs
TYPE: nucleic acid
STRANDEDNESS: Sirc
TOPOLOGY:
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OTHER INFORMATION: PCR Primer US-09-213-768-2
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US-08-259-451
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Best Local
Matches 1
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Patent No. 64000
Patent INFORMATION:
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TYPE: DNA
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COMPUTER
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
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MEDIUM TY
COMPUTER:
OPERATING
                                                                                                                                                                                                         FILING DATE: 20
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                         COUNTRY
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TYPE: Diskette
TYPE: storage
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5406841
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Abbott Park
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                                                                                                                                                                                                                                                                                                                                                        Illinois
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Johnson, Joan E.
Motley, Cheryl T.
Peterson, Bryan
Edwards, Michelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Golde, David W. Robertson, Euger Stephens, John E Chan, Emerson W.
                                                                                                                                                                                                                                                                                                                                          USA
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Conservative
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Tate, Cynthia
                                                                                                                                                                                                                                                                                                                                                                                         Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESS:
                                                                                                                                                                                                                     20-JUN-1994
                                                                                                                                                                                                                                                                                           storage
                                                                                                                                                                                                                                                                                                       Diskette,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Emerson W.
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Pred. No.
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US-08-259-451-10

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ADDRESSUL
STREET: P. O. BUA
CITY: Houston
STREET: P. O. BUA
CITY: HOUSTON
STATE: TX
COUNTRY: USA
ZIP: 77210

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249,380
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/07/710,187
FILING DATE: 31-MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: ARCD:024
TELEPHONE: 713-787-1400
TELEPHONE: 713-787-1400
TELEFAX: 713-789-2679
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 3727 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-249-380-1

US-08-249-380-1

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Search completed: July Job time: 49.2525 secs
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US-08-249-380-1/c
; Sequence 1, Application US/08249380
; Patent No. 5827685
; GENERAL INFORMATION:
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Best Local S
Matches 18
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CORRESPONDENCE ADDRESS: APPLICATION White & STREET: P. O. Box 4433
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TITLE OF IN
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Response Systems
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Pred. No. 19;
0; Mismatches
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19;
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version - 2003

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Title:
Perfect so
Sequence:
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Maximum
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cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result		% Query Match	Length	BB	ID	Description
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14		•	9		0-146-726-31	319, A
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## ALIGNMENTS

RESULT 1

US-10-010-408-1

Sequence 1, Application US/10010408

Publication No. US20020165185A1

Publication No. US20020165185A1el Heparin-Induced CCN-Like Molecules

APPLICANT: John J. Castellot, Jr.

ADDRESSE: LAHIVE & COCKFIELD, LLP

STREET: 28 State Street

CITY: Boston

STATE: Massachusetts

CONFUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER READABLE PORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENIAN Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: US-DOS/MS-DOS

FILING DATE: OT-Dec-2001

CLASSIFICATION NUMBER: US-DOS/MS-DOS

FILING DATE: OT-Dec-2001

PRIOR APPLICATION UNMBER: US-DOS/MS-DOS

FILING DATE: OT-Dec-2001

FRICATON DATE: MARCH 19, 1998

APPLICATION UNMBER: 35,207

RESISTRATION UNMBER: 35,207

RESISTRATION UNMBER: MSI-004

TELEFORMULICATION LOWER: MSI-004

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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
FEATURE:
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LOCATION: 249..10
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
RENT APPLICATION DATA:
                                                                                                                                                                                                                                                 NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-
ECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
ION FOR SEQ ID NO: 3:
JENCE CHARACTERISTICS:
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       NAME/KEY:
                                                                                                                                                                                                            TOPOLOGY:
                                                                      GTGTGTGCCCAGCTGTGCCGGACACCCTGTACCTGTCCTTGGACACCACCCCAGTGCCCA
                                                CAGGGGGTACCCCTGGTGCTGGATGGCTGTGGCTGCTGTAAAGTGTGTGCACGGAGGCTG
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APPLICATION NUMBER: 09/044,27
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
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                                                                                                     imilarity 100.0%;
; Conservative
                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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                                        AGGGGGTACCCCTGGTGCTGGATGGCTGTGGCTGCTGTAAAGTGTGTGCACGGAGGCTG
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RESULT 3
US-10-010-408-12
; Sequence 12, Application US/10010408
; Publication No. US20020165185A1
; GENERAL INFORMATION:
: APPLICANT: John J. Castellot, J.
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                                                                                                             INFORMATION
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FILING DATE: 07-Dec-2001
CLASSIFICATION LATA:
APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
REGISTRATION NUMBER: 36,207
REGISTRATION NUMBER: 36,207
                                                                      REFERENCE/DOCKET NUMBER: 36,207
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
ORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 681 bacc
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COMPUTER READABLE
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
CULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
STREET: 28
CITY: Bosto
STATE: Mass
COUNTRY: US
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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                                                          TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Boston
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Y: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           State Street
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US-10-010-408
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Best Local S
Matches 566
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                                                                                      WBER OF SEQUENCES: 13
RRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & CONTY: 28 State Streets
CITY: Boston
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
MPUTER READABLE FORM:
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LOCATION: 1..681
UENCE DESCRIPTION: S
8-12
                                                                                                                                                                                                                                                       ICANT: John J. Castellot, Jr.
E OF INVENTION: No. US20020165185Alel Heparin-Induced CCN-Like
and Uses Therefor
                                                                                                                                                                                                                                                                                                               NFORMATION:
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
ENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                Application US/10010408 No. US20020165185A1
                                                                                                                                                                                                                                                                                                                                                                                                                                           CAGCTGTGCCGGACACCCTGTACCTGTCCTTGGACACCACCCCAGTGCCCACAGGGGGGTA
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ilarity 100.0%;
Conservative
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Pred. No. 1.3e-284;
0; Mismatches 0;
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                       #1.0,
                       Version
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NAME/KEY: CDS;
LOCATION: 1..210;
SEQUENCE DESCRIPTION: 105-10-010-408-8
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US-10-010-408-5
                                                                                                                                                                                                                                         Sequence 5, Application US/10010408
Publication No. US20020165185A1
GENERAL INFORMATION:
APPLICANT: John J. Castellot
TITLE OF INVENTION: No. US20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 23.8%;
Best Local Similarity 100.0%;
Matches 210; Conservative
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                        ZIP: 02109

ZIP: 02109

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, V

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)742-4214
ORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                   NUMBER OF SEQ
CORRESPONDENC
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ADDRESSEE: LAHIVE & COCSTREET: 28 State Street
CITY: Boston
STATE: Massachusetts
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APPLICATION NUMBER: FILING DATE: 07-Dec-
                                                                                                              COUNTRY: USA
ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                            CCTGGCGGCCATGGGGCTGTGTGTCTCTTG
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                                                                                                                                                                                                                 SEQUENCES:
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and Uses Therefor
S: 13
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Pred. No. 3.9e-9
0; Mismatches
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US/10/010,408
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APPLICANT: COhen, Robert
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Levine, Arnold J.
APPLICANT: Levine, Arnold J.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC ACIDS ENC
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/10/112,267
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-04
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-04
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-14
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-14
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US-10-112-267-
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US-10-010-408-5
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Best Local
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TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
ORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 177 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match
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STRANDEDNESS: single
TOPOLOGY: linear
CULE TYPE: cDN2
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OR APPLICATION DATA:
APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
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CATION: 1..177
E DESCRIPTION: S
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larity 100.0%;
Conservative
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RESULT 8
US-10-010-408-10
; Sequence 10, Application US/10010408
; Publication No. US20020165185A1
; GENERAL INFORMATION:
APPLICANT: John J. Castellot, J.
TITLE OF INVENTION: No. US20020
and Uses The
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CURRENT APPLICATION NUMBER: US/10/112,267
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR FILING DATE: 1998
PRIOR FILING DATE: EARLIER FILING DATE: 1997
PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR FILING DATE: 1998
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ORGANISM: Mus r
S-10-112-267-17
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TYPE: DNA
ORGANISM: M
S-10-112-267-1
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Best Local
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Similarity 100.0%;
90; Conservative
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Goddard,
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SEQUENCES:
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, William I.
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ce, David A.
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              Castellot, Jr.
No. US20020165185A1el
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                                                                                                                                                                                                                                                                                                                              Score 90;
Pred. No
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DATE: 1998-10-29
R APPLICATION NUMBER:
DATE: 1997-10-29
R APPLICATION NUMBER:
DATE: 1998-02-04
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Mismatches
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hes 0;
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1.1e-36;
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                               Heparin-Induced
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                               CCN-Like
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                                Molecules
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NAME/KEY: CDS;
; LOCATION: 1..17;
; SEQUENCE DESCRIPTION:
US-10-010-408-10
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US-09-864-761
      APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME
TITLE OF INVENTION: GENE EXPRES
FILE REFERENCE: Aeomica-X-1
CURRENT APPLICATION NUMBER: US/0
CURRENT FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/
PRIOR APPLICATION NUMBER: US 60/
PRIOR APPLICATION NUMBER: US 60/
PRIOR APPLICATION NUMBER: US 09/
PRIOR APPLICATION NUMBER: US 09/
PRIOR APPLICATION NUMBER: GB 242
PRIOR APPLICATION NUMBER: GB 242
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                        Sequence 2 Patent No.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
ORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 174 ---
                                                                                                                                                                                ICANT: Chen, Wensheng
E OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC
E OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
REFERENCE: Aeomica-X-1
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COMPUTER READABLE FORM:
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ADDRESSEE: LAHIVE
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FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
IOR APPLICATION DATA:
APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
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STRANDEDNESS: single
TOPOLOGY: linear
CULE TYPE: cDNA
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COUNTRY: US
ZIP: 02109
ICATION NUMBER:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Amy E. Mandragouras REGISTRATION NUMBER: 36,2
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Rank, David R.
Hanzel, David K.
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llarity 100.0%;
Conservative
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60/236,359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MBI-004
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1.8e-20;
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Sequence 6698, Application US/09864; Patent No. US20020048763A1; GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-D
TITLE OF INVENTION: GENE EXPRESSI
FILE REFERENCE: Aeomica-X-1; CURRENT APPLICATION NUMBER: US/09/
CURRENT FILING DATE: 2001-05-23; PRIOR APPLICATION NUMBER: US 60/18
PRIOR APPLICATION NUMBER: US 60/20
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Best Loc
Matches
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TYPE: DNA
ORGANISM:
FEATURE:
OTHER INFO
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364-761-6698
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OR APPLICATION NUMBER: PCT/US01/00665
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00668
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00663
OR APPLICATION NUMBER: PCT/US01/00663
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 60/234,687
OR APPLICATION NUMBER: US 60/234,687
OR APPLICATION NUMBER: US 09/608,408
OR APPLICATION NUMBER: US 09/608,408
OR FILING DATE: 2000-06-30
OR APPLICATION NUMBER: US 09/774,203
OR FILING DATE: 2001-01-29
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APPLICATION NUMBER: PCT/US01/00669
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EXPRESSED IN PLACENTA, SIGNAL = 1.

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EXPRESSED IN FETAL LIVER, SIGNA

EXPRESSED IN BONE MARROW, SIGNA

EXPRESSED IN ADULT LIVER, SIGNA

EXPRESSED IN HELA, SIGNAL = 2.7

EXPRESSED IN BRAIN, SIGNAL = 1.

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NUMBER: PCT/
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TYPE: DNA
ORGANISM: HOMO Sapid
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OTHER INFORMATION: NOTHER INFORMA
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DR APPLICATION NUMBER: US 09/632,366

DR FILING DATE: 2000-08-03

DR APPLICATION NUMBER: GB 24263.6

DR FILING DATE: 2000-10-04

DR APPLICATION NUMBER: US 60/236,359

DR APPLICATION NUMBER: PCT/US01/00666

DR FILING DATE: 2001-01-30

DR APPLICATION NUMBER: PCT/US01/00667

DR APPLICATION NUMBER: PCT/US01/00667

DR APPLICATION NUMBER: PCT/US01/00669

DR APPLICATION NUMBER: PCT/US01/00669

DR APPLICATION NUMBER: PCT/US01/00665

DR APPLICATION NUMBER: PCT/US01/00668

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DR APPLICATION NUMBER: PCT/US01/00668

DR APPLICATION NUMBER: PCT/US01/00663

DR APPLICATION NUMBER: PCT/US01/00662

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DR APPLICATION NUMBER: US 60/234,687

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                                                                                                                                                                                                                  INFO
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INFORMATION:
-761-6698
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                                                Goddard, Audrey
Gurney, Austin L.
Hillan, Kenneth J.
Lawrence, David A.
                                                                                                                                                                                       Botstein,
                                                                                                                                                                                                                                              Application US/10112267 No. US20030068678A1
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N HEART, SIGNAL = 1.9

N FETAL LIVER, SIGNAL = 1

N BONE MARROW, SIGNAL = 1

N ADULT LIVER, SIGNAL = 1

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US-10-112-267-39
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CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION
PRIOR FILING DATE: EARLIER FILING DATE: 1990
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APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDIFILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/10/11
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER AIPPRIOR FILING DATE: EARLIER FILING DIPPRIOR FILING DATE: EARLIER FILING D
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EQ ID NO 38
LENGTH: 738
TYPE: DNA
ORGANISM: Homo
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LENGTH: 84
TYPE: DNA
ORGANISM:
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NAME/KEY:
LOCATION:
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Goddard, Audrey
Gurney, Austin
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Pennica, Diane
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e, David A.
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DATE: 1998-10-29
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APPLICATION NUMBER:
DATE: 1998-04-14
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
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PR APPLICATION NUMBER: 60/089947

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PR APPLICATION NUMBER: 60/089947

PR FILING DATE: 1998-06-24

PR APPLICATION NUMBER: 60/090449

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RESULT 14 US-10-146-7 ; Sequence Qy Query Mato Best Local Matches 36 75  $\omega N$ 16 3.6%; Similarity 100.0%; 2; Conservative 0 20 319 Score 32; DB 14; Pred. No. 1.8e-06; 0; Mismatches 0; Length 1266; 0 Gaps

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Application US/10146726

APPLICANT: APPLICANT: ENERAL

INFORMATION:

Baker, Kevin

Beresini,

Maureen

APPLICANT:

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APPLICANT:
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                                           APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C312
CURRENT APPLICATION NUMBER: US/10/146,727
CURRENT FILING DATE: 2002-05-15
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http://image.llnl.gov  Plate: LLAM14017 row: n column: 21  High quality sequence stop: 543.  Location/Qualifiers  1940	Email: cgapbs-remail.nin.gov Tissue Procurement: The Cepko Laboratory cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can be found through the I M A G F Consortium/ILNI.at.	Contact: Robert Strausberg, Ph.D.	Unpublished (1999)	National Institutes of Health, Mammalian Gene Collection (MGC)	NIH-MGC http://mgc.nci.nih.gov/.	1 to 940)	Rodentia;	Riikarvota: Metagoa: Chordata: Craniata: Vertehrata: Riiteleoetomi:	Mile mile Cilline	house mouse.	EST.	BQ937887.1 GI:22353365	BQ937887	nRNA sequence	AGENCOURT_8951807 NCI_CGAP_Co24 Mus musculus cDNA clone	BQ937887 940 bp mRNA linear EST 21-AUG-2002	

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Assembly, verification, and initial annotation of NIA 7.4K mc cDNA clone set
Unpublished (2002)
Contact: Yong Qian
Laboratory of Genetics
National Institute on Aging/National Institutes of Health
333 Cassell Drive, Suite 3000, Baltimore, MD 21224-6820, USA
Email: cdna@lgsun.grc.nia.nih.gov
This clone set has been freely distributed to the community.
visit http://lgsun.grc.nia.nih.gov/cDNA/NIA_7_4k.html for det
Plate: H4067 row: A column: 01
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/clone=lib="NCI_CGAP_Co24"
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/note="Organ: colon; Vector: p@MV-SPORT6; Site_1:
/note="Organ: cloned unidirectionally. Primer: Oleverage insert size 1.6 kb. Constructed by Life
Average insert size 1.6 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
9 a 277 c 288 g 200 t 6 others
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/db_xref="taxon:10090"
/clone="H4067A01"
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/lab_host="DH10B"
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/sex="mixed"
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                                                                                                                                                 Watahiki, M., ..., S., Kawai, J., Okazaki, Y., Muramarsu, ...,
Hayashizaki, Y.
RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
Konno, H., Fukunishi, Y., Shibata, K., Itoh, M., Carninci, P., Sugahara, Y. and Hayashizaki, Y.
Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) please visit our web site (http://genome.gsc.riken.go.jp) for
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S Akimura, T., Arakawa, T., Carninci, P., Furuno, M., Hanagaki, T.,

Hayatsu, N., Hiramoto, K., Hiraoka, T., Hirozane, T., Imotani, K., Ishii
, Y., Ito, M., Kawai, J., Kojima, Y., Konno, H., Kouda, M., Matsuyama, T.,

Nakamura, M., Nishi, K., Nomura, K., Numasaki, R., Okazaki, Y., Okido, T.,

Saito, R., Sakai, C., Sakai, K., Sakazume, N., Sasaki, D., Sato, K.,

Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Suzuki, H., Tagawa, A., Takahashi, F., Takaku-Akahira, S., Tanaka, T., Tomaru, A., Toya, T.,

Watahiki, A., Yasunishi, A., Muramatsu, M. and Hayashizaki, Y.,

RIKEN Encyclopedia of Mouse Full-length cDNAs (Akimura, T., et al. 2001)

L Unpublished (2001)

Contact: Yoshihide Hayashizaki
Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), Yokohama Institute

The Institute of Physical and Chemical Research (RIKEN)
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Fax:
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URL:http://genome.gsc.riken.go.jp/
Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K.,
,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.

Normalization and subtraction of cap-trapper-selected cDNAs t
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. 10 (10), 1617-1630 (2000)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1-7-22 Suehiro-cho,
Tel: 81-45-503-9222
Fax: 81-45-503-9216
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BB849097 RIKEN full-length enriched, adu cDNA clone F930006G02 5', mRNA sequence.
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house mouse.
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e Institute of Physical and Chemical Research (RIKEN)
7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Jaj
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3.8e-20;
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inner (
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      adult
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BG900020
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HOA48-1-G2.R HOA
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Eukaryota; Me
Mammalia; Eut
1 (bases 1 t
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BG900020.1
EST.
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Identification and initial characterization sequenced tags (ESTs) each from adult human osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001) 21482651
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                                                                                                 32;
                                                                                                                                                                                                                                                                                                                         Tel:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Le
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Similarity 100.0%;
49; Conservative
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                                                GTACCCCTGGTGCTGGATGGCTGTGGCTGCTG
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L: 610-270-7245
k: 610-270-5598
                                                                                                                                                                                                                                                                                     primer:
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                                                                                                                                                                                                                                                                                     sanjay_kumar-1@gsk.com
.mer: T7.
                                                                                                                                                           Sanjay Kumar
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/dev_stage="adult"
/note="pooled tissues; (tissue_type=cerebellum,
    dev_stage=16 days neonate, sex=mixed),
    (tissue_type=cerebellum, dev_stage=0 day neonate,
    (tissue_type=hippocampus, dev_stage=adult,
    sex=male), (tissue_type=whole body, dev_stage=9 days
    embryo, sex=mixed), (tissue_type=lung, dev_stage=13 day
    embryo, sex=mixed)"
    124 c 132 g 103 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Metazoa;
Eutheria;
1 to 380)
                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="HOA (Human Osteoarthritic
/tissue_type="cartilage"
/lab_host="E.coli DH10 B"
                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                               organism-"Homo
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Primates;
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Pred. No. 7.9e-12;
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BM751866
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BG900069

HOA51-1-All.R HOA (Human Osteoarthritic Cartilage) Hom cDNA, mRNA sequence.

BG900069

BG900069.1 GI:14310318

EST.

human.

M Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu Mammalia; Eutheria; Primates; Catarrhini; Hominidae; H 1 (bases 1 to 405)

Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., G Lark,M.W.

Identification and initial characterization of 5000 ex sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries

Osteoarthr. Cartil. 9 (7), 641-653 (2001)

Contact: Sanjay Kumar

UW2109

GlaxosSmithKline

709 Swedeland Road, P.O. Box 1539, King of Prussia, PA Tel: 610-270-7245

Fax: 610-270-5598

Email: sanjay_kumar-1@gsk.com

Seq primer: T7.
BM751866

K-EST0028180 S2SNU668s1 Hom
5', mRNA sequence.
BM751866
BM751866.1 GI:19081484
EST.
human.
Homo sapiens
Eukaryota; Metazoa; Chordat
Mammalia; Eutheria; Primate
1 (bases 1 to 489)
Kim, N.S., Hahn, Y., Oh, J.H.,
Oh, K.J., Cheong, J.E., Sohn,
Kim, Y.S.
21C Frontier Korean EST Pro
Unpublished (2002)
Contact: Kim YS
Genome Research Institute of
52 Eoeun-dong Yuseong-gu, I
Tel: +82-42-860-4470
Fax: +82-42-860-4409
Email: yongsung@mail.kribb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   375
                                                                                                                                                                                                                                                                                                                                    M751866 489
-EST0028180 S2SNU668s1 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 100.0%;
2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                    GTACCCCTGGTGCTGGATGGCTGTGGCTGCTG 406
      yongsung@mail.kribb.re.kr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pSPORT I;
Directional"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /oryanism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/tissue_type="cartilage"
/lab_host="E.coli DH10 B"
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1. .405
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Primates; Catarrhini;
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Pred. No.
                                                                                                                                                          J.H., Lee,J.Y.,
Sohn,H.Y., Kim,
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                                                of Bioscience & B
, Daejeon 305-333,
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J.M., Park,H.S., Kim,S. and
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                                                   Biotechnology
3, South Korea
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S2SNU668s1-2-D09
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Matches 32
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euto
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 
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mRNA sequence.
BG538695
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/db_xref="taxon:9606"
/clone="S2SNU668s1-2-D09"
/clone_lib="S2SNU668s1"
/sex="m"
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/cell_line="SNU-668"
/lab_host="Topl0F'"
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IMAGE:4691574 5',
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BG928868
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BG928868.;
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GlaxoSm
709 Swe
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Mammalia; E
                                                                                                                                                                                                                                                          Identification and initial characterization sequenced tags (ESTs) each from adult human osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001) 21482651 Contact: Sanjay Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                               human
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larity 100.0%;
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/db_xref="taxon:9606"
/clone=lib="NIH_MGC_77"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
Sfil (ggccgctcggcc); Site_2: Sfil (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
5'-ATTCTAGAGGCCGAGGCCGACATG-dT(30)BN-3' (where B = A,
C; or G and N = A, C, G, or T). Average insert size 1.9
kb (range 0.5-4.0 kb). 12/15 colonies contained inserts
by PCR. This library was enriched for full-length clones
and was constructed by Clontech Laboratories (Palo Alto,
CA). Note: this is a NIH_MGC Library."

95 a 194 c 213 g 116 t
                                                                                                                                                             sanjay_kumar-l@gsk.com
imer: T7.
                        /db_xref="taxon:9606"
/clone_lib="HNC (Human Normal C./tissue_type="cartilage"
/lab_host="E.coli DH10 B"
/note="Vector: pSPORT I; Site_1
Directional"
               Ø
                                                                                                                                                                                                                                                                                                                                                                                                      ; Metazoa;
Eutheria;
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                                                                                                                organism="Homo
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i; Hominidae;
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NIH-MGC http://mgc.no
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BI457141
BI457141.1 GI
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prime, mRNP
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Similarity 100.0%;
32; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
Unpublished (1999)
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Location/Qualifiers
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                                                                                                                                                                        /image.llnl.gov
LLCM1869 row:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       cgapbs-r@mail.nih.gov
Procurement: ATCC
                    mRNA
                                                                                                                                                                       Hong In California, Be (Stratagene) / Note: this is 230 c
                              LTI_NFL006_PL2
                                                                                                                                                                                                                                                                                                                                          organism="Homo sapiens
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 GI:12896595
                    sequence
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Primates;
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Catarrhini
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IMAGE:5258159 5',
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ON 603620978F1 NIH_MGC_40 Homo sapiens cDNA clone image.

NRNA sequence.
NRNA sequence.
NRNA sequence.
NRNA sequence.
NRM43988.
BM043988.
IGI:16773255

EST.
ISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 750)
RS NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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Similarity 100.0%;
2; Conservative
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                                                                                                                                        quality sequence
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                                                                                                  ity sequence stop:
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/note="Vector: pCMVSPOR
                                         ′organism="Homo
xref="taxon:9606"
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Pred. No.
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RESULT 12
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Matches 32
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 790)

S NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1944 row: h column: 17
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603626068F1 NIH_MGC_40 Homo s
mRNA sequence.
BM046275
BM046275.1 GI:16775542
EST.
3.6%;
Similarity 100.0%;
32; Conservative
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Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   quality sequence stop:
Location/Qualifiers
                                                                                            /Organism="Homo sapiens"
/db_xref="taxon:9606"
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Site_2: EcoRI; cDNA made by oligo-dT priming.
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI, XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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/tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oldgo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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IMAGE:5452480
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E 1 (bases 1 to 800)

S NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

L Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA Library Preparation: Life Technologies, Inc.

CDNA Library Preparation: Life Technologies, Inc.

Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL)

Plate: LIAM11420 row: g column: 18
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                                                                                                                                                                                                                         375
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BI822142
603039845F1 NII
mRNA sequence.
BI822142
BI822142.1 GI
EST.
human.
Homo sapiens
Eukaryota; Meta
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603077268F1 NII
mRNA sequence.
BI826781
BI826781.1 GI
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human.
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3077268F1 NIH_MGC_119
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                                                                                                                                                                                                                                                                                                                                               Site_2: EcoRV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note this is a NTU NCC Titure.
                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Organ: brain; Vector:
Site_2: EcoRV (destroyed); R)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /uryanism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="medulla"
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/note="Orange
    Metazoa;
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IMAGE: 5169065 5
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    Euteleostomi;
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Eukaryota; Metazoa; (
Mammalia; Eutheria; |
1 (bases 1 to 888)
NIH-MGC http://mgc.n.
National Institutes
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32; Conser
                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution informat
found through the I.M.A.G.E. Consortium/LLNL at:
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Plate: LLAM11450 row: j column: 11
                                   http://image.llnl.
Plate: LLAM11408
                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
Unpublished (1999)
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Location/Qualifiers
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Tissue Procurement: Life Technologies,
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llarity 100.0%;
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pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA
source anonymous pool of 6 male brains, age range 23-27; 1
male lung, age 27; and 1 male testis, age 69. Library is
oligo-dT primed and directionally cloned (EcoRV site is
destroyed upon cloning). Average insert size 1.8 kb,
insert size range 1-3 kb. Library is normalized and
enriched for full-length clones and was constructed by C.
Gruber (Invitrogen). Research Genetics tracking code
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/db_xref="taxon:9606"
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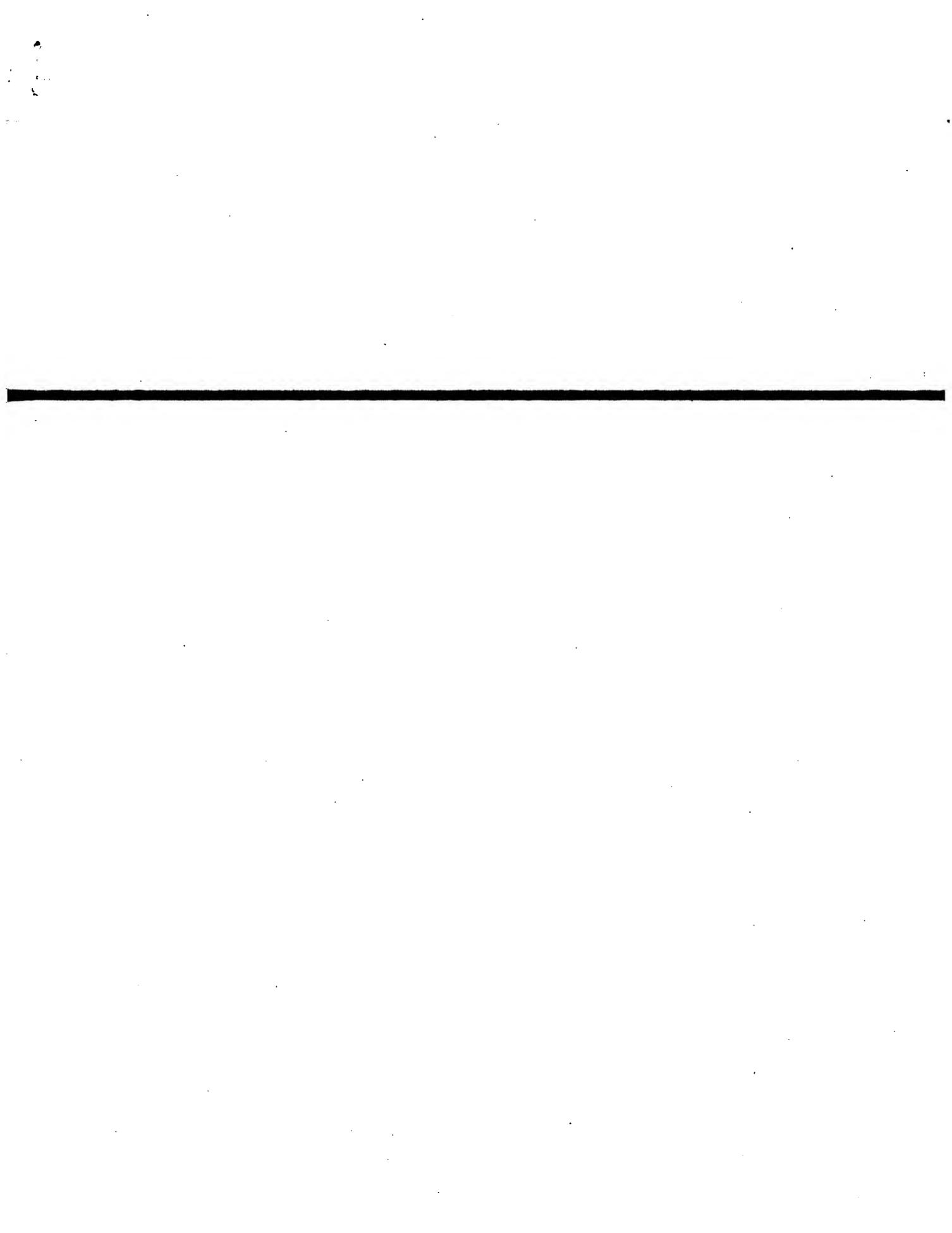
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llarity 100.0%;
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/clone_lib="NIH_MGC_119"
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/lab_host="DH10B"
/note="Organ: brain; Vectors
                                                                                                                                                                                                  /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note: this is a NIH_MGC Library."
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REFERENCE AUTHORS		SOURCE ORGANISM	KEYWORDS	VERSION	ACCESSION	DEFINITION	RESULT 1 AF259981 LOCUS	
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ted (24-APR-2000) Cell Biology, Vangerbilt-
, 649 MRB II, Nashville, TN 37232, USA
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/product="CCN family protein COP-1"
/product="CCN family protein COP-1"
/protein_id="AAF69011.1"
/db_xref="GI:7739781"
/translation="MRGSPLIRLLATSFLCLLSMVCAQLCRTPCTCPWTPPQCPQGVPLVLDGCGCCKVCARRLTESCEHLHVCEPSQGLVCQPGAGPGGHGAVCLLDEDDGDCEVNGRRYLDGETFKPNCRVLCRCDDGGFTCLPLCSEDVTLPSWDCPRPKRIQVPGKCCPEWVCDQGVTPAIQRSAAQGHQLSALVTPASADAPWPNWSTAWGPCSTTCGLGIATRVSNQNRFCQLEIQRRLCLPRPCLAARSHSSWNSAF"
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/gene="Cop-1"
/note="secreted
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/db_xref="taxon:10116"
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Direct Submission
Submitted (24-JUL-2002) Human
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Baylor Plaza, Houston, TX 7703
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1 (bases 1 to 137964)

1 (bases 1 to 137964)

Muzny,D.M., Adams,C., Adio-Oduola,B., Ali-osman,F.R., Alieu, Alisbrooks,S.L., Amaratunge,H.C., Are,J.R., Ayele,M., Banks,T. Alisbrooks,S.L., Benton,J., Bimage,K., Blankenburg,K., Bonnin,D.
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Direct Submission
Submitted (10-JUL-2002) Human Genome Sequencing Center, Depair of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA

3 (bases 1 to 137964)
       * 1
                                                     Center project name: GZHG
Center clone name: CH230-301E4
Center clone name: CH230-301E4
Center clone name: CH230-301E4
Center clone name: CH230-301E4
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 89474 bases at least Q40
Consensus quality: 93422 bases at least Q30
Consensus quality: 936506 bases at least Q20
 NOTE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mitted (24-JUL-2002) Human Genome Sequencing Center, Depa
Molecular and Human Genetics, Baylor College of Medicine,
lor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                  Web site: http://www.tmc.edu
Contact: hgsc-help@bcm.tmc.edu
----- Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Center: Baylor College of Medicine
Center code: BCM
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Cleveland, C.D., Cox, C., Coyle, M.D., Datnorne, S.K., David, R., Davis, C., Davy-Carcoll, L., Déderich, D.A., Delandy, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delunthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Edwards, C.C., Elhaß, C., Escotto, M., Esc
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2 (bases 1 to 226303)
Worley, K.C.
Direct Submission
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3 (bases 1 to 226303)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (11-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor college Baylor Plaza, Houston, TX 77030, USA On Jul 10, 2002 this sequence version replaced contermination.
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 Assembly program: Phrap; version Consensus quality: 185950 bases a Consensus quality: 190362 bases a Consensus quality: 193076 bases a
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Estimated insert size may differ from sequence 10 e http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_0 This is a 'working draft' sequence. It currently sts of 49 contigs. The true order of the pieces t known and their order in this sequence record is rary. Gaps between the contigs are represented as of N, but the exact sizes of the gaps are unknown record will be updated with the finished sequence on as it is available and the accession number wi
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1 (bases 1 to 1734)

Botstein, D.A., Cohen, R.L., Goddard, A.D., Gurney, A. Lawrence, D.A., Levine, A.J., Pennica, D., Roy, M.Ann. WISP polypeptides and nucleic acids encoding same Patent: US 6387657-A 18 14-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                             Unknown
Unknown
                                                                                                                                                                                                                                                              90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unclassified.
1 (bases 1 to 1734)
1 (bases 1 to 1734)
Botstein,D.A., Cohen,R.L., Goddard,A.D., Gurney,A.L., Hillan,K:J
Botstein,D.A., Levine,A.J., Pennica,D., Roy,M.Ann. and Wood,W.I.
Lawrence,D.A., Levine,A.J., Pennica,D., Roy,M.Ann. and Wood,W.I.
WISP polypeptides and nucleic acids encoding same
Patent: US 6387657-A 17 14-MAY-2002;
Location/Qualifiers
1. .1734
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sh 12.0%;
Similarity 100.0%;
90; Conservative
                                                                         Mus musculus connective (Wisp2) mRNA, complete of AF100778
AF100778.1 GI:4028578
                                                                                                                     AF100778
                                                                                                                                                                                                                                                            Similarity 100.
90; Conservative
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(bases 1 nnica, D.,
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 Swanson, T.A.,
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                    Chordata;
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tissue g
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US 6387657.
Welsh, J.W.,
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                 Craniata; Ver
Sciurognathi;
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Roy,M.A.,
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                            Vertebrata;
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                    Muridae;
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Lawrence, D.
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                               Euteleostomi;
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                    Murinae;
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>tein WISP-2
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SOURCE
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Best Local
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sequence.

AL731698
AL731698.10 GI:21214309
HTG.
house mouse.

Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
E 1 (bases 1 to 61072)
S Wallis, J.
Direct Submission
Submitted (24-MAY-2002) Wellcome Trust Sanger Institute, Hinxton,
Cambridgeshire, CB10 1SA, UK. E-mail enquiries:
humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On May 25, 2002 this sequence version replaced gi:21213601.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lee, J., Brush, J., Taneyhill, L.A., Deuel, B., Lew, M., Watanabe, C., Cohen, R.L., Melhem, M.F., Finley, G.G., Quirke, P., Goddard, A.D., Hillan, K.J., Gurney, A.L., Botstein, D. and Levine, A.J. WISP genes are members of the connective tissue growth factor family that are up-regulated in wnt-1-transformed cells and aberrantly expressed in human colon tumors

L. Proc. Natl. Acad. Sci. U.S.A. 95 (25), 14717-14722 (1998)

E. 99061933

D. 9843955

E. 2 (bases 1 to 1734)

S. Pennica, D.

Direct Submission

L. Submitted (23-OCT-1998) Molecular Oncology, Genentech, Inc., 1 DNA Way, South San Francisco, CA 94080, USA
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Mouse DNA
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1. .1734
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WVCDQAVMQPAIQPSSAQGHQLSALVTPASADGPCPNWSTAWGPCSTTCGLGIATRVS
NQNRFCQLEIQRRLCLSRPCLASRSHGSWNSAF"
491 c 495 g 393 t
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/codon_start=1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /gene="Wisp2"
257. .1012
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                                                                                                                                                                                                                                                                                                                                             sequence from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /cell_line="C57MG"
/cell_type="epithelial"
/tissue_type="mammary"
/note="transformed by Wnt-1"
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Pred. No.
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1.8e-37;
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                                                                                               Sims, S.

Sims, S.

Direct Submission

L Submitted (23-JUL-2002) Wellcome Trust Sanger Institute, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail enquiries:
humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On Jul 25, 2002 this sequence version repliaced gi:18181793.

Center: Wellcome Trust Sanger Institute
Center code: SC

The otter http://www.sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         222
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  Chemistry:
                                                                                                                                                                                                                                                                                                                             Eukaryota;
Mammalia; E
                                only a small overlap as described above.

This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers give in the feature table with their source databases: Em:, EMBL; Sw:, SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at
                                                                                    Web site: http://www.sanger.ac.uk
Contact: humquery@sanger.ac.uk
----- Project Information
                                                                                                                                                                                                                                                                                                                                                                                                  AL669906
AL669906.5 GI:21955520
HTG; HTGS_PHASE1; HTGS_0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     AL669906 216757 bp DNA Mus musculus chromosome 2 clone RP23-217C2, PROGRESS ***, 21 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://www.sanger.ac.uk/Projects/C_elegans/wormpep RP23-161B from the RPCI-23 Mouse PAC Library constructed by the group of Pieter de Jong. For further details see http://www.chori.org/bacpac/home.htm
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a 15368 c 15033 g
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/db_xref="taxon:10090"
  Dye-terminator quality: 211865
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Eutheria; Rodentia;
1 to 216757)
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Pred. No.
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Mismatches 0;
Big Dye;
bases at
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Sciurognathe; Muridae; Murinae; Mus
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least Q40
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100 by 136 84635: contig of 1
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10 87499: cr
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46360: contig of 6589 bp in length

61 46460: gap of 100 bp

61 49809: contig of 3349 bp in length

10 49909: gap of 100 bp

10 54509: contig of 100 bp
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                                                                                                       vector
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3 15367: σ
                                                                                                                                      note="assembly_fragment:03429
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                                                                                                                                                                                                                                                  'organism="Mus musculus"
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5 213877: conti
6 213977: gap of
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39671: c
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216757: cont
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                                               AGTGTGTGCACGGAGGCTGGGGGAGTCCTGCGACCACCTGCATGTCTGCGACCCCAGCCA
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54610. .6
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                                                                                                                                              Score 90; DB Pred. No. 9.5 0; Mismatches
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0358067
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0; Conservative
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Biol. Chem. 274 (24), 17123-17131 (1999)
 R210337
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                                                                                                                                                                                                                                                        /protein_id="AAD18058.1"
/db_xref="GI:4337060"
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WVCDQAVMQPAIQPSSAQGHQLSALVTPASADGPCPNWSTAWGPCSTTCGLGIATRVS
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480 c 489 g 395 t
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/gene="Ctgfl"
/note="similar to the Mus musculus WISP-2 protein encoded
/note="similar to the Mus musculus WISP-2 protein encoded
by the sequence presented in GenBank Accession Number
AF100778; putative growth factor; CTGF-L; contains IGF
binding (IGFBD), Von Willebrand Factor type C (VWC) repeat
and thrombospondin type I (TSP1) domains; member of the
CCN (CTGF/Cyr61/Nov) family; lacks the fourth
carboxy-terminal (CT) domain present in other members of
the CCN family"
                                                                                                                                                                                                                                                                                                                                                                      precursor"
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1. .1739
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242. .997
                                                                                                                                                                                                                                                                                                                                                                                     /codon_start=1
/product="connective tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Mus musculus"/db_xref="taxon:10090"
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|. .1739
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Eukaryota; Metazoa; C
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AR210337:
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1 (bases 1 to 738)
Botstein, D.A., Cohen, R.L., Goddard, A.D., Gurney, A.L.
Botstein, D.A., Levine, A.J., Pennica, D., Roy, M.Ann. a
Lawrence, D.A., Levine, A.J., Pennica, D., Roy, M.Ann. a
WISP polypeptides and nucleic acids encoding same
Patent: US 6387657-A 38 14-MAY-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                1 (bases 1 to 841)
Botstein, D.A., Cohen, R.L., Goddard, A.D., Gurney, A.L.,
Lawrence, D.A., Levine, A.J., Pennica, D., Roy, M.Ann. and
WISP polypeptides and nucleic acids encoding same
Patent: US 6387657-A 39 14-MAY-2002;
Location/Qualifiers
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32; Conservative
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Unclassified.
Polypeptidic compositions
Patent: WO 0105836-A 31 2
                    Botstein,D.,
Wood,W.I.
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larity 100.08;
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Primates;
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25-JAN-2001;
                              Gurney, A.L.,
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Catarrhini;
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8.4e-06;
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d Wood,W
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AUTHORS
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Homo sapiens.

SM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 1283)

S Kumar,S., Hand,A.T., Connor,J.C., Dodds,R.A., Ryan,P.J.,

Trill,J.G., Fisher,S.M., Slemmon,J.R., Lipshutz,D.B.,

Bartholomew,V., James,I.E., Rieman,D.J., Gowen,M. and Lee,J.C.

Identification and cloning of CTGF-L from human osteoblasts, a novel cysteine rich protein containing an IGF binding domain

Bone 23 (5), S240 (1998)

E 2 (bases 1 to 1283)
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Secreted and transmembrane polypeptides and nucleic acids encodin same
Patent: WO 0140466-A 319 07-JUN-2001;
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/db_xref="taxon:9606"
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/db_xref="taxon:9606"
418 c 390 g 24
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8e-06;
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Direct Submission
Submitted (11-AUG-1998) Bone & Cartilage Biology, UW 2109,
SmithKline Beecham, 709 Swedeland Road, King of Prussia, PA 19406,
USA
                                                                                                                                        th 4.2%; S
Similarity 100.0%;
32; Conservative 0;
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July 28,
                                                                                                                                                                                                                          /note="CTGF-L; encodes IGF binding (IGFB), Von Willebrand factor type C (VWC) and thrombospondin type I (TSP1) domains; member of the CCN (CTGF/cyr61/nov) family; lacks the fourth carboxy-terminal domain present in other members of the CCN family"
/codon_start=1
/product="connective tissue growth factor-like protein /precursor"
/protein_id="AAC70350.1"
/protein_id="AAC70350.1"
/translation="MRGTPKTHLLAFSLLCLLSKVRTQLCPTPCTCPWPPPRCPLGVPLVLDGCGCCRVCARRLGEPCDQLHVCDASQGLVCQPGAGPGGRGALCLLAEDDSSCEVNGRLYREGETFQPHCSIRCRCEDGGFTCVPLCSEDVRLPSWDCPHPRRVEVLGKCCPEWVCGQGGGLGTQPLPAQGPQFSGLVSSLPPGVPCPEWSTAWGPCSTTCGLGMATRVSNQNRFCRLETQRRLCLSRPCPPSRGRSPQNSAF"
35 a 418 c 389 g 241 t
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/db_xref="taxon:9606"
/chromosome="20"
/map="20q12-q13"
/cell_type="primary osteoblast"
9. .761
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1. .1283
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                  21:14:49
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O; Mismatches
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## ALIGNMENTS

26-NOV-1999

(first entry)

polypeptide coding sequence.

standard; cDNA; 753 BP.

induced CCN-like protein; HICP; cell-associated activity; ss; scular disorder; aberrant cell proliferation; fibrotic disorder.

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     acid sequences encoding rat heparin-induced CCN-like protein, methods to identify modulators or in diagnostic applications
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RESULT 3
AAZ07521
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AC AAZ
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AC AAZ
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DE Rat
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CCA
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PF 18-
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       ention provides a rat heparin-induced CCN-like protein (HICP). Agents that stimulate or inhibit HICP protein activity or ion, antisense HICP nucleic acid molecules and HICP antibodies, used to modulate cell-associated activity. HICP modulators can be treat disorders characterized by aberrant HICP protein activity ession. Probes capable of hybridizing to HICP mRNA or antibodies c for HICP can be used to detect HICP activity in a biological
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RESULT 4
AAZ07519
ID AAZ07519
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AC AAZ07519;
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DT 26-NOV-19
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DE Rat HICP
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KW Heparin-i
KW cardiovas
KW insulin-l

99

standard;

cDNA;

210

IGFBP

encoding

cDNA.

induced CCN-like protein; HICP; scular disorder; aberrant cell like growth factor binding prot

HICP; cell-associated activity; cell proliferation; fibrotic di protein; IGFBP.

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Similarity 100.0%;
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                        CTTGTCACTCCTGCCTCTGCTGATGCTCCTTGTCCAAATTGGAGCACAGCCTGGGGCCCC
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Pred. No. 2.4e-312;
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RESULT 5
AAZ07518
ID AAZ0
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AC AAZ0
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DT 26-N
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DE Rat
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KW Hepa
KW Card
KW Von
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Matches 210; Conserva
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P-PSDB;
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                     Heparin-induced CCN-like cardiovascular disorder; von Willebrand C domain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used
                                                                                                        AA207518;
 Rattus
                                                                                                                            AA207518
                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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DB; AAY27438.
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Pred. No. 2.1e-89;
D; Mismatches 0;
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                              cell-associated activity; ss;
proliferation; fibrotic disorder;
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P-PSDB; AA
  18-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA; 174 BP.
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                                                                                                                                                                                    protein; HICP; cell-associated activity; ss; aberrant cell proliferation; fibrotic disorder; TSP1.
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RESULT 7
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AC AAX7
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DT 06-2
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CONT
KW WNT-
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P-PSDB;
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                                                                                                                       connective tissue growth factor; cancer; melanoma; arteriosclerosis leukaemia; lymphoid malignancy; haematopoiesis-related disorder; tissue-growth disorder; skin disorder; desmoplasia; fibrotic lesion kidney disorder; bone-related disorder; osteoporosis; trauma; burn; connective tissue disorder; catabolic state; inflammation; testicular-related disorder; angiogenesis; immunological disorder;
                                                                                                                                                                                                                                                06-AUG-1999
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  14-APR-1998;
                                                  06-MAY-1999
                                                                          W09921998-A1.
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                          29-OCT-1998;
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Pred. No. 2.3e-7
0; Mismatches
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                                                                                                                                                                                                 WISP-1;
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lators or in diagnostic appl
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e-72;
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RESULT 8
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KW Conne
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Cantibodies can be used to induce death in WISP-1, 2 or 3 overexpressing
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Best Local S
Matches 90
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997;
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                                                                                                                                                                                                                                                                                        standard; DNA;
                                                                                                                                                                                                                                                                                                                                                            GGGCCTGGTTTGTCAGCCTGGGGCAGGCCC
                                                                                                                                                                                                                                                                                                                                                                                        GGGCCTGGTTTGTCAGCCTGGGGCAGGCCC 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 753 BP; 159 A; 229 C; 238 G; 127 T;
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larity 100.0%; Pred. No. 1.3e-32;
Conservative 0; Mismatches 0;
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Levine AJ
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98US-0073612.
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kidney disorder; bone-related disorder; osteoporosis; trau connective tissue disorder; catabolic state; inflammation; testicular-related disorder; angiogenesis; immunological of

disorder; ss.

WO992199

8-A1.

999.

gp

connective tissue growth factor; cancer; melanoma; arteriosclerosis; leukaemia; lymphoid malignancy; haematopoiesis-related disorder; tissue-growth disorder; skin disorder; desmoplasia; fibrotic lesion; kidney disorder; bone-related disorder; osteoporosis; trauma; burn;

induced secreted protein; WISP-1; WISP-2; WISP-3; CTGF;

WISP-2 protein nucleotide sequence SEQ ID NO:17.

WNT-1

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RESULT 9
ABA72245
ID ABA7
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AC ABA7
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DT 01-F
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DE Huma
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Antibodies can be used to induce death in WISP-1, 2 or 3 overexpressing
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Best Local S
Matches 90
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29-OCT-1997;
03-FEB-1998;
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P-PSDB;
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Lawrence
                                                      Homo
                                 WO200157277-A2
                                                                                                                     01-FEB-2002
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DB; AAY17651.
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                                                                                               foetal
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Similarity 100
90; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Page 178-179;
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Levine
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97US-0063704.
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RESULT 10
ABA38112
ID ABA38112
XX
AC ABA38112;
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DT 23-JAN-20
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DT 23-JAN-20
XX
Cardiovas
KW Human; ge
KW congenita
XX
OS Homo sapi
XX
PN WO2001572
XX
PD 09-AUG-20
XX
PF 30-JAN-20
PR 04-FEB-20
PR 30-JUN-20
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Note: The s
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                              Human; gene expression; cardiovascular disease; congenital heart disease
                                                                                                                               30-JAN-2001;
                                                                                                                                                                                WO20015727
                                                                                                                                                                                                                                                                                                                                    ABA38112;
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                                                                                                                                                                                                                                                                                   78
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                                                                                                                                                                                                                                                                                                                                                                                                                           99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence data for this patent did not form part of the ecification, but was obtained in electronic format directly at ftp.wipo.int/pub/published_pct_sequences.
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    LECULAR
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                                                                                                                                                                                                                                  heart disease; h
                           2000US-0180312.

2000US-0207456.

2000US-0608408.

2000US-0632366.

2000US-0234687.

2000US-0236359.

2000US-0236359.
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larity 100.0%;
Conservative
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2000US-0207456.

2000US-0608408.

2000US-0632366.

2000US-0234687.

2000US-0236359.

2000US-024263.
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                                                                                                                                2001WO-US00666
                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO 20550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32 A;
                                                                                                                                                                                                                                                                                  gene expression analysis
    DYNAMICS
                                                                                                                                                                                                                                                                                                                                                              DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65 C; 73 G;
                                                                                                                                                                                                                                   hypertension;
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                                                                                                                                                                                                                                                            heart;
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3.8e-05;
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RESULT 11
AAK20667
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AC AAK20
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AC AAK20
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O5-NC
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O5-NC
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Humar
KW Humar
KW micrc
KW micrc
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Best Local S
Matches 32
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                Single
brains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples defrom the human heart via microarrays. By measuring gene expression, probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular e.g. cardiovascular disease, hypertension, cardiac arrhythmias and congenital heart disease.

Note: The sequence data for this patent did not form part of the puspecification, but was obtained in electronic format directly from at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                microarray; Alzhe epilepsy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-2001
                                                                                                                                                                                                                                                                                                 30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human;
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2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000GB-0024263.
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                   present invention provides a number of single exon nucleic acid es which are derived from genomic sequences expressed in the human marrow. They can be used to measure gene expression in bone marrow les, which may enable the improved diagnosis and treatment of cancers as lymphoma, leukaemia and myeloma. The present sequence is one of probes of the invention.
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Similarity 100.0%;
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cancer; leukaemia; lymj
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2000US-0632366.

2000US-0234687.

2000US-0236359.

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                                                                                                                                                                                                                                                (SENP). The present sequence is one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printe specification, but was obtained in electronic format directly from WIPC at ftp.wipo.int/pub/published_pct_sequences.
                              AAİ52650
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connected acid expressed in the human lung; measuring gene expression in a sample derived from human lung, comprising (a) contacting the array with a collection of detectably labeled nucleic acids derived from human lung measuring gene expression in a collection of detectably labeled nucleic acids derived from human lung manual acids derived from human lung manual acids derived from human lung manual acids derived from human lung manual acids from expression in a eukaryotic genome, comprising charles acids from expression and post detectably bound to each probe of the expression and (b) detecting at least one exon from genomic sequences of the eukaryote; and (b) detectably leaven genome, comprising comprising a fragment identical to the predicted exon, the probe is included in the above mentioned microarray; assigning exons to a single exon probe, having a fragment identifying exons from genomic sequence by the method above and (b) measuring the expression of each of the xons in several tissues and/or cell types using hybridisation to a single gene, comprising one the exons should be assigned to a single gene, particularly using human lung derived mRNA the probes are used for gene expression analysis, and for identifying exons in a gene, particularly using human lung derived mRNA and for the study of lung disease (COPD), interstitial lung disease (ILD), familial idiopathic pulmonary fibrosis, pulmonary histiocytosis, lymphangiolelomyomtosis, pulmonary bysplasia, primary ciliary dyskinesis, pulmonary hypertension and hyaline membrane disease. The present sequence is a single exon probe open reading frame of the invention.

To fibrosi the formation this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614 probes. Also included are a microarray comprising the novel set of probes; the novel set of probes which hybridise at high stringency to
                                                                 Sequence
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-145-18/c = 18, Application US/09182145B NO. 6387657 INFORMATION:

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APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND N
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
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APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NI
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER FILING DATE: 1998-02-04
EARLIER FILING DATE: 1998-04-14
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APPLICANT: Levine, Arnold J.
APPLICANT: Pennica, Diane
APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUC
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT FILING DATE: 1998-10-29
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CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
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                                                                                                                                                                         Sequence
Patent No
GENERAL J
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Sequence 13,
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                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 39
LENGTH: 841
TYPE: DNA
ORGANISM: Artificial
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ICANT: Wood, William I.
3 OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC REFERENCE: P1176R2
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Gurney, Austin L.
Hillan, Kenneth J.
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Gurney, Austin L.
Hillan, Kenneth J.
Lawrence, David A.
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GENERAL INFORMATION:
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CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Levine, Arnold J.
APPLICANT: Pennica, Diane
APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P1176R2
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EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
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TYPE: DNA
ORGANISM: HOMO
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ORGANISM: H
09-182-145-1
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APPLICANT:
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LENGTH: 1293
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APPLICATION NUMBER:
FILING DATE: 1998-0
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32; Conservative
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Goddard,
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      Audrey
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P., David A.
        Austin
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; TYPE: DNA
; ORGANISM: H
; FEATURE:
; HAME/KEY: m
; LOCATION: (
; OTHER INFOR
US-09-385-982-2
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TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION

TITLE OF INVENTION: PRODUCTS: II

FILE REFERENCE: CCDNA-260XX

CURRENT APPLICATION NUMBER: US/09/385,982

CURRENT FILING DATE: 1999-08-30

EARLIER APPLICATION NUMBER: 09/328,111

EARLIER FILING DATE: 1999-06-08

EARLIER APPLICATION NUMBER: 60/117,393

EARLIER APPLICATION NUMBER: 60/098,639

EARLIER APPLICATION NUMBER: 60/098,639

EARLIER FILING DATE: 1998-08-31
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CURRENT FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1997-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
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LENGTH: 61
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LENGTH: 51
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9-182-145-117
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INVENTION: WISP POLYPEPTIDES AND NUCLEIC ACIDS ENCODING SAME SRENCE: P1176R2
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Sequence 225, Applica sequence 225, Applica sequence 225, Applica sequence 225, Applica sequence 225, Applica sequence 225, Applica sequence 226, Application 6420526

GEARLIER FILING DATE 11 Sequence 220 CURRENT APPLICATION 220 CURRENT APPLICATION 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DATE 220 CURRENT FILING DAT
ANT: ROSED et al.

WEFERENCE: PZ002P1

WEFERENCE: PZ002P1

YT APPLICATION UNMBER: US/09/149,47,47

YT FILING DATE: 1998-09-08

RAPPLICATION NUMBER: 60/040,162

ER APPLICATION NUMBER: 60/040,666

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/040,33

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

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ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,50

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PLICATION NUMBER: 60/047,593
LING DATE: 1997-05-23
PLICATION NUMBER: 60/047,614
PLICATION NUMBER: 60/043,578
POTICATION NUMBER: 60/043,578
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PLICATION NUMBER: 60/043,578
LING DATE: 1997-04-11
LING DATE: 1997-04-11
PLICATION NUMBER: 60/043,576
LING DATE: 1997-04-11
LING DATE: 1997-05-23
PLICATION NUMBER: 60/047,501
LING DATE: 1997-04-11
PLICATION NUMBER: 60/056,632
LING DATE: 1997-08-22
PLICATION NUMBER: 60/056,632
LING DATE: 1997-08-22
PLICATION NUMBER: 60/056,876
LING DATE: 1997-08-22
PLICATION NUMBER: 60/056,876
LING DATE: 1997-08-22
PLICATION NUMBER: 60/056,881
LING DATE: 1997-08-22
PLICATION NUMBER: 60/056,881
LING DATE: 1997-08-22
PPLICATION NUMBER: 60/056,887
PLICATION NUMBER: 60/056,884
PLICATION NUMBER: 60/056,884
PLICATION NUMBER: 60/056,884
PLICATION NUMBER: 60/056,884
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Similarity 100.0%;
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LICATION NUMBER: 60/05
PLICATION NUMBER: 60/04
TNG DATE: 1997-06-13
                                                                                                                                                                                                                                                                                                                                TGATGACGGTGGCTTCACC
                                                                                               Brenda F. Baker
Lex M. Cowsert
VENTION: ANTISENSE MODULATION OF
NCE: RTS-0026
                                   ICATION NUMBER: US/09/213,768
ING DATE: 1998-12-17
EQ ID NOS: 47
                                                                                                                                                                                                     opplication
085664
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ICATION NUMBER: 60/
ING DATE: 1997-10-02
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NG DATE: 1997-05-23
ICATION NUMBER: 60/047,590
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ICATION NUMBER:
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NUMBER: 60/
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US-
                                   Sequence 2, Application US/09213768; Patent No. 5985664; GENERAL INFORMATION:
APPLICANT: Brenda F. Baker; APPLICANT: Lex M. Cowsert; TITLE OF INVENTION: ANTISENSE MODULA FILE REFERENCE: RTS-0026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-
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US-09-213-768-2/c
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FILE REFERENCE: 790CIP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
NUMBER OF SECTION NUMBER: 09/540,217
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Best Local S
Matches 19
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APPLICANT: Drmanac, Radoj
TITLE OF INVENTION: No. (
TITLE OF INVENTION: Polyp
FILE REFERENCE: 790CIP2A
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LOCATION: (136
-09-213-768-1
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         CURRENT APPLICATION NUMBER: US/09/213,768
CURRENT FILING DATE: 1998-12-17
                                                                                                                                                                                                                                                                                                                                         LENGTH: 1539
TYPE: DNA
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
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NUMBER OF
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LOCATION: (130)..(1539)
)9-668-680-13
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Local Similarity 100.0%;
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Zhang, Jie
Wang, Jian-Rui
Xue, Aidong J.
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Liu, Chenghua
Zhou, Ping
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Sequence Patent No

No.

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2, Application US/09182145B 6387657

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RESULT 15
US-09-182-145
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US-08-249-380
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Best Local :
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 0
FILING DATE: 31-MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lindquist, Susan
TITLE OF INVENTION: Methods and Compositions
TITLE OF INVENTION: Stress Response Systems
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-787-1400
TELEFAX: 713-789-2679
                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0'
FILING DATE: 31-MAY-1991
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CITY:
STATE:
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R READABLE FORM:
M TYPE: Floppy disk
TER: IBM PC compatible
TING SYSTEM: PC-DOS/MS-DOS
NARE: PatentIn Release #1.0, V
APPLICATION DATA:
NAPPLICATION DATA:
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Pred. No. 19;
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APPLICANT: Cohen, Robert
APPLICANT: Goddard, Audrey
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Lawrence, David A.
APPLICANT: Levine, Arnold J.
APPLICANT: Levine, Arnold J.
APPLICANT: Pennica, Diane
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUC:
FILE REFERENCE: P1170R2
CURRENT APPLICATION NUMBER: US/09/182,145B
CURRENT APPLICATION NUMBER: US 60/063,704
EARLIER APPLICATION NUMBER: US 60/063,704
EARLIER FILING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: US 60/073,612
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: US 60/081,695
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 156
SEQ ID NO 152
LENGTH: 44
TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: 1-44
OTHER INFORMATION: Sequence is synthesized.
Patent No. 6387657
US-09-182-145-152
Search completed: July 28, Job time: 42.1485 secs
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Matches 17
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APPLICANT: Cohen, Robert
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Similarity 100.0%;
17; Conservative
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Perfect score:
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1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

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8: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

10: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

11: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

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16: /cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*

17: /cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*
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Sequence 3, Application US/10010408
Publication No. US20020165185A1

GENERAL IMPORMATION:

APPLICANT: John J. Castellot, Jr.

APPLICANT: John J. Castellot, Jr.

APPLICANT: John J. Castellot, Jr.

TITLE OF INVENTION: No. US20020165185A1el Heparin-Induced CCN-Like Molecules

ITLE G DATE: DATE: LAHIVE & COCKFIELD, LLP

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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
FEATURE:
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LOCATION: 1..750
SEQUENCE DESCRIPTION: :
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COMPUTER READABLE FORM:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
RENT APPLICATION DATA:
APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
OR APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
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NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
ECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
ION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
CULE TYPE: cDNA
GAGGTGAATGGCCGCAGGTACCTGGATGGAGAGACCTTTAAACCCAATTGCAGGGTCCTG
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LOCATION: 249..1001
NCE DESCRIPTION: SEQ
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ADDRESSEE: LAHIVE
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OF INVENTION: No. US20020165185Alel Heparin-Induced CCN-Like Molecules
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RESULT 3
US-10-010-408-12
: Sequence 12, Application US/10010408
; Publication No. US20020165185A1
; GENERAL INFORMATION:
: APPLICANT: John J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J. Castellot, J.
                                                                                                TELEFAX: (
                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vei
CURRENT APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
TELEPHONE: (617)227-7400
TELEPHONE: (617)742-4214
FORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 681 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
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COMPUTER READABLE FORM:
MEDIUM TYPE: Flopp
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STATE: Massac
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ADDRESSEE: LAHIVE
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Publication N
GENERAL INF
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Best Local
Matches 68
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LOCATION: 1..681
SEQUENCE DESCRIPTION: 0-408-12
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ADDRESSEE: 1
STREET: 28:
CITY: Boston
STATE: Mass:
                                                                                                                                           Application US/10010408 No. US20020165185A1 FORMATION:
                                               CANT: John J. Castellot, Jr.
OF INVENTION: No. US20020165185Alel
and Uses Therefor
R OF SEQUENCES: 13
SPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD, LLP
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US-10-010-408-8
                                                                                                                                             Sequence 5, Application Publication No. US2002 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 27.9
Best Local Similarity 100.
Matches 210; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Verent Application DATA:
APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatibl

COMPUTER: IBM PC-DOS/1
                                                              NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE
                                                                                                                                   APPLICANT:
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CITY: Boston
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: CDS
LOCATION: 1..210
NCE DESCRIPTION:
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                                                   STREET: 28
                                                                                                                    INVENTION:
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: No. US20020165185A1el
and Uses Therefor
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tIn Release #1.0,
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Pred. No.
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                                                                                                                                                                                                                                                                  sequence 10, Application Publication No. US20020 GENERAL INFORMATION:

APPLICANT: Tohn
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Matches
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APPLICATION NUMBER: US/10/010,408
FILING DATE: 07-Dec-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: 09/044,273
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: March 19, 1998
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: Number: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)742-4214
VFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 177 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity nes 177; Conserv
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, 1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match
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CORRES
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FEATURI
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CITY: Boston
STATE: Massac
                                                                                                 STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                      ANT: John J. Castellot, Jr.
OF INVENTION: No. US20020165185A1el Heparin-Induced CCN-Like
and Uses Therefor
OF SEQUENCES: 13
PONDENCE ADDRESS:
ADDRESSE: LAHIVE & COCKFIELD, LLP
TREET: 28 State Street
                                                                                                                                                                                                                                                                                                          Application US/10010408 to. US20020165185A1
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TYPE: nucleic acid
TRANDEDNESS: single
TOPOLOGY: linear
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larity 100.0%;
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Pred. No. 9.1e-82;
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                    APPLICANT: ROY, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AN
FILE REFERENCE: P1176R2
CURRENT APPLICATION NUMBER: US/10/112,26
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLIC
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ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: MBI-004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)742-4214
PORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 174 base pairs
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TOPOLOGY: linear
MOLECULE TYPE: cDNA
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Similarity 100.0%;
74; Conservative
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APPLICATION NUMBER: <Unknown>
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Goddard, Audrey
Gurney, Austin
Hillan, Kenneth
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Pennica, Diane
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             DATE:
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R APPLICATION NUMBER: 1997-10-29
            APPLICATION NUMBER: DATE: 1998-04-14
                                    APPLICATION DATE: 1998-0
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RESULT 9 US-09-864-761

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Sequence

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NO.

432, Application US20020048763A1

US/09864761

GENERAL IN APPLICANT

INFORMATION:

Penn,

Sharron G David R.

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CURRENT APPLICATION NUMBER: US/10/112,267
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/PRIOR FILING DATE: EARLIER FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/PRIOR FILING DATE: EARLIER FILING DATE: 1997-10-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-04
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TYPE: DNA
ORGANISM:
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Wood, William I.
NVENTION: WISP POLYPEPTIDES AND NUCLEIC ACIDS ENCODING
ENCE: Pl176R2
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Gurney, Austin
Hillan, Kenneth
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Lawrence, David A.
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                                                                                                        Score 90; DB 15; L; Pred. No. 1.1e-36;
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TYPE: DNA
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PR APPLICATION NUMBER: PCT/US01/00665

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PR APPLICATION NUMBER: PCT/US01/00668

PR APPLICATION NUMBER: PCT/US01/00668

PR APPLICATION NUMBER: PCT/US01/00663

PR APPLICATION NUMBER: PCT/US01/00662

PR APPLICATION NUMBER: PCT/US01/00661

PR APPLICATION NUMBER: PCT/US01/00661

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PR APPLICATION NUMBER: US 60/234,687

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R FILING DATE: 2001-01-30
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R APPLICATION NUMBER: US 09
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R APPLICATION NUMBER: GB 24
R FILING DATE: 2000-10-04
R APPLICATION NUMBER: US 60
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IN PLACENTA, SIGNAL -
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083500.1, EVALUE 1
HIT: 019113, EVALU
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PRAPPLICATION NUMBER: PCT/US01/00666

PRILING DATE: 2001-01-30

PRAPPLICATION NUMBER: PCT/US01/00664

PRILING DATE: 2001-01-30

PRAPPLICATION NUMBER: PCT/US01/00669

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PRAPPLICATION NUMBER: PCT/US01/00668

PREILING DATE: 2001-01-30

PRAPPLICATION NUMBER: PCT/US01/00663

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                       APPLICANT: Roy, Margaret Ann
APPLICANT: Wood, William I.
TITLE OF INVENTION: WISP POLYPEPTIDES AND NUTTITLE OF INVENTION: WISP POLYPEPTIDES AND NUTTITLE OF INVENTION NUMBER: US/10/112,267
CURRENT APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR APPLICATION NUMBER: EARLIER APPLICATION PRIOR FILING DATE: 1991
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LENGTH: 738
TYPE: DNA
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5 DATE: 1997-10-29
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DATE: 1998-02-04
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FILE REFERENCE: P3330R1C151
CURRENT APPLICATION NUMBER: US/10/137,866
CURRENT FILING DATE: 2002-05-07
PRIOR APPLICATION:
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NUMBER: 60/C

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R APPLICATION NUMBER: 60/059115
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R APPLICATION NUMBER: 60/059122
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R APPLICATION NUMBER: 60/059352
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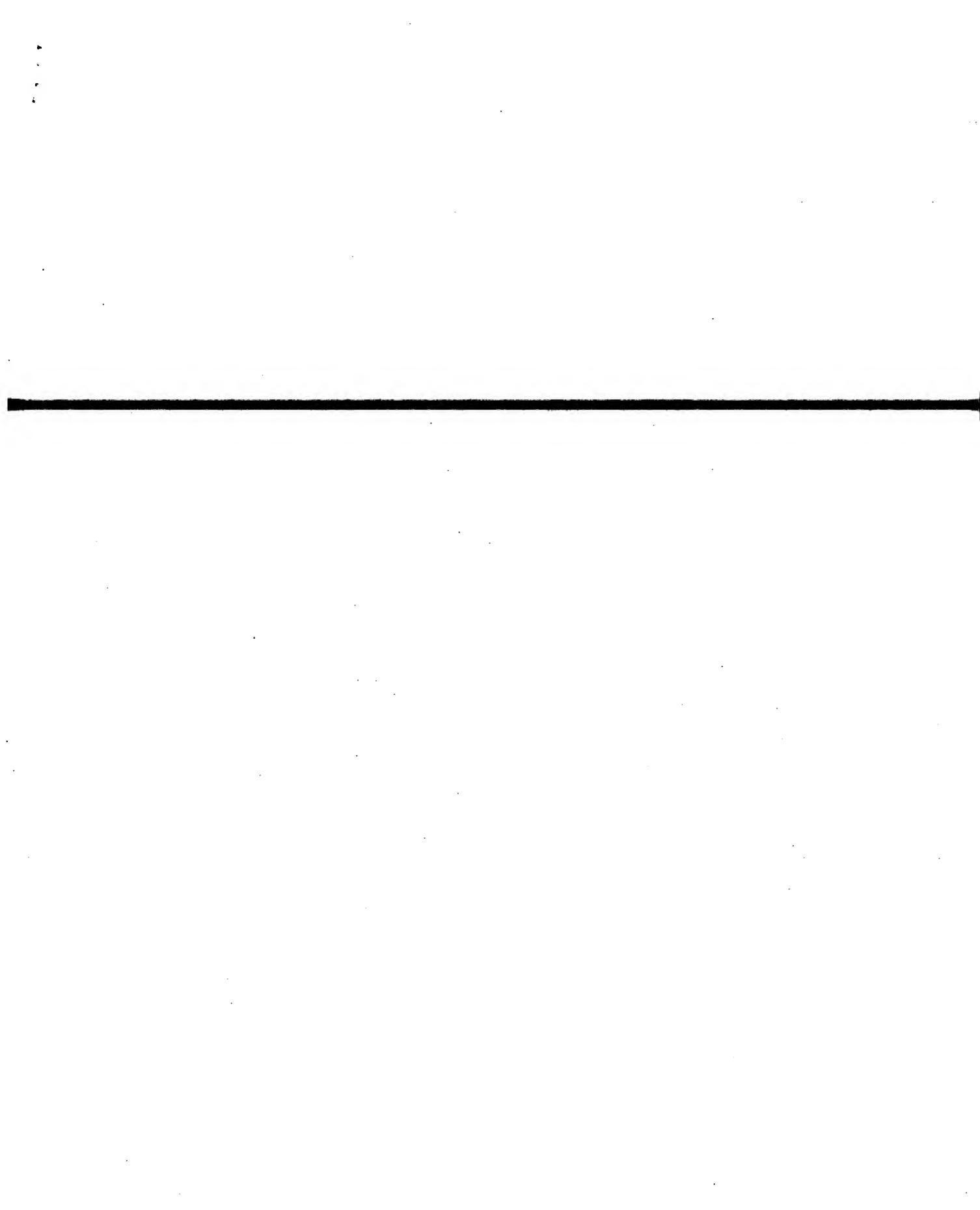
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C308
CURRENT APPLICATION NUMBER: US/10/146,726
CURRENT FILING DATE: 2002-05-15
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 319
LENGTH: 1266
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Gao, Wei-Qiang
Gerritsen, Mary E.
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Stewart, Timothy A.
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o. US20030129690A1
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FILE REFERENCE: P3330R1C312
CURRENT APPLICATION NUMBER: US/10/146,727
CURRENT FILING DATE: 2002-05-15
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 550
SEQ ID NO 319
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## ALIGNMENTS

Definition AGENCOURT\_8951807 NCI\_CGAP\_CO24 Mus musculus cDNA clone

IMAGE:6475852 5', mRNA sequence.

ACCESSION B937887
VERSION EST.
SOURCE DAYS HOUSE Musculus
EXTRORDS EST.
ORGANISM Mus musculus
ELKaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
AUTHORS NATIONAL Institutes of Health, Mammallan Gene Collection (MGC)
COMMENT Email: cgapbs-r@mail.nih.gov/
Tissue Procurement: The Cepko Laboratory
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.lnl.gov
light procurements of the Stribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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Laboratory of Genetics
National Institute on Aging/National Institutes of Health
333 Cassell Drive, Suite 3000, Baltimore, MD 21224-6820, Email: cdna@lgsun.grc.nia.nih.gov
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. Similarity 100.0%;
90; Conservative
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This clone set has been freely distributed to visit http://lgsun.grc.nia.nih.gov/cDNA/NIA.plate: H4067 row: A column: 01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VanBuren, V., Piao, Y., Dudekula, D.B., Qian, Y., Cart, P.R., Stagg, C.A., Bassey, U., Aiba, K., Hamatani, T. Luo, A.G. and Ko, M.S.H. Assembly, verification, and initial annotation of
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/note="Organ: colon; Vector: pCMV-
                  /note-"Vector: pSPORT1; clone is among a rearra; than 20 cDNA libraries. 162 c 166 g 1
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/strain="EVB/N"
/db_xref="taxon:10090"
/clone="IMAGE:6476852"
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/strain="C57BL/6"
                                                                                                                                                               /clone_lib="NIA Mouse
/sex="mixed"
                                                                                                                   /dev_stage="mixed"
/lab_host="DH10B"
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/db_xref="taxon:10090"
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Best Local S
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                                                                                                                                                                                                                                                  Watahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and Hayashizaki,Y.

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Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a encyclopedia: real-time sequence clustering for construction of a
                                                                                                                                                                                                                                                                                                                                                                                                                           Email: genome-res@gsc.riken.go.jp,

URL:http://genome.gsc.riken.go.jp/

Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K.,
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genes. Genome Res. 10 (10), 1617-1630 (2000)

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,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and
Hayashizaki,Y.
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Please visit
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BB849097 RIKEN full-length enriched, adu cDNA clone F930006G02 5', mRNA sequence.
BB849097
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ntact: Yoshihide Hayashizaki
boratory for Genome Exploration Research Group, RIKEN
lences Center(GSC), Yokohama Institute
e Institute of Physical and Chemical Research (RIKEN)
7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-
1: 81-45-503-9222
x: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             imura,T., Arakawa,T., Carninci,P., Furuno,M., Hanagaki,T.,
yatsu,N., Hiramoto,K., Hiraoka,T., Hirozane,T., Imotani,K., Ishii
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              caryota;
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                                                           /strain="Mus musculus"
/strain="C57BL/6J"
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                     /clone="F930006G02"
                                               /db_xref="taxon:10090"
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                                                                                                                                                                                                               sed methods for the mouse full-length cDNA: real-time sequence clustering for construction of cDNA library. Genome Res. . 11 (2), 281-289 (2001) tour web site (http://genome.gsc.riken.go.jp) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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Pred. No. 3.9e-22;
0; Mismatches 0;
full-length enriched,
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ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                         Waterston, R.
The WashU-HHMI Mouse EST Project
Unpublished (1996)
Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Loui
Tel: 314 286 1800
Fax: 314 286 1810
Email: mouseest@watson.wustl.edu
This clone is available royalty-free through
IMAGE Consortium (info@image.llnl.gov) for i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                373
                                                                                                                                                                                                                                                                                                                                                   MGI:642454
Seq primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA754979 488 bp mRNA
vu55g08.rl Soares_mammary_gland_NbMMG Mus
IMAGE:1195358 5', mRNA sequence.
AA754979
AA754979.1 GI:2802177
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               49;
                                                                                                                                                                                                                                                                                                                 Seq primer: -28ml3 rev2 ET from High quality sequence stop: 478 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrick
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Mc
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K.,
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson
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49; Conservative
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(tissue_type=cerebellum, dev_stage=0 day neonate,
sex=mixed), (tissue_type=hippocampus, dev_stage=adult,
sex=male), (tissue_type=whole body, dev_stage=9 days
embryo, sex=mixed), (tissue_type=lung, dev_stage=13 day
embryo, sex=mixed)"

124 c 132 g 103 t
                               adaptors
                                                                                 primer
                                                                 1. .488
                                                                                 /note-"Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco
RI; 1st strand cDNA was primed with a Not I - oligo(dT)
primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eutheria;
                                                                                                                                                                                                                                 /organism="Mus musculus
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:1195358"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="inner ear"
/dev_stage="adult"
/note="pooled tissues;
                                                                                                                                                 /tissue_type="mammary
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                  /clone_lib="Soares_mammary_gland_NbMMG
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 to 488)
 provided
                 Not
double-stranded cDNA was ligated to Eco RI (Pharmacia), digested with Not I and cloned I and Eco RI sites of the modified pT7T3 vect vided by Dr. Minoru Ko, Wayne State Univ. Libra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 49; DB. Pred. No. 3.1 0; Mismatches
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Sciurognathi;
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hi; Muridae;
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COMMENT

**FEATURES** 

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ACCESSION
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BF141695

BF141695.1 GI:10980735

EST.

house mouse.

SM Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.

1 (bases 1 to 966)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Gilbert Smith, Ph.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by:Incyte Genomics, Inc.

Clone distribution: NCI-CGAP clone distribution information can 1

found through the I.M.A.G.E. Consortium/LLNL at:
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http://image.llnl.gov
Plate: LLAM9276 row: h
High quality sequence:
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601790752F1 NCI_CGAP_Lu30 Mus musculus
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Similarity 100.0%;
35; Conservative
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mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/strain="CZECH II"
/db_xref="taxon:10090"
/clone="IMAGE:4021308"
/clone_lib="NCI_CGAP_Lu30"
/tissue_type="tumor, metasta/lab_host="DH10B"
                                                                                                                                                                                                                                                  /note="Organ: lung; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: SalI; transgenic model WNT-1, expression driven MMTV-LTR enhancer; Cloned unidirectionally. Primer: OlidT. Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH" 298 c 258 g 183 t
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                                                                                                                                                                           red. No. 3e-06;
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                                            380 bp
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Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W.,
Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee
Lark,M.W.
Identification and initial characterization of
sequenced tags (ESTs) each from adult human no
osteoarthritic cartilage cDNA libraries
Osteoarthr. Cartil. 9 (7), 641-653 (2001)
21482651
Contact: Sanjay Kumar
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Tel: 610-270-7245
Fax: 610-270-5598
Email: sanjay_kumar-1@gsk.co
Seq primer: T7.
                                  GlaxoSmithKline
709 Swedeland Road,
Tel: 610-270-7245
Fax: 610-270-5598
                                                                                                                                                                                                                                                                                                                                                Homo sapiens
Eukaryota; Me
Mammalia; Eut
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709 Swedeland Ro
                                                                                                                                                        sequenced tags (ESTs) each from adult human osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001) 21482651
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Sathe,G.,
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Identification and initial
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 sanjay_kumar-1
.mer: T7.
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/db_xref="taxon:9606"

/clone_lib="HOA (Human Osteoarthritic Calissue_type="cartilage"

/lab_host="E.coli DH10 B"

/note="Vector: pSPORT I; Site_1: Salt. coli principle of the coli principle of the coli principle of the coli principle of the coli principle of the coli principle of the coli principle of the coli principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin principle of the colin princi
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                                                                                                                                         Sanjay Kumar
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Eutheria;
1 to 405)
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### Eutheria; Primates; Catarrhini; F
1 to 380)
Connor, J.R., Dodds, R.A., Halsey, F
Mui, P., Agarwal, P., Badger, A.M.,
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Mui, P., Agar
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Lee,J.C.
                                                                           Prussia,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 489)
Kim, N.S., Hahn, Y., Oh, J.H., Lee, J
Oh, K.J., Cheong, J.E., Sohn, H.Y.,
Kim, Y.S.
21C Frontier Korean EST Project 2
Unpublished (2002)
Contact: Kim YS
Genome Research Center
Korea Research Institute of Biosc
52 Eoeun-dong Yuseong-gu, Daejeon
Tel: +82-42-860-4470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             202
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K-ESTO
5', mR
BM7518
BM7518
EST.
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Tel: +82-42-860-4470
Fax: +82-42-860-4409
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|751866
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Location/Qualifiers
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/note="Organ: Stomach; Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly (A)+ RNA was decapped with tabacco acid pyrophosphatase (TAP) and ligated with DNA-RNA linker including EcoRI site by treatment of T4 RNA ligase. The first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli Top10F' by electroporation method. After analyzing and sequencing about 2,000 ~ 3,000 colonies in original cDNA library, the abundant cDNAs were
                                                                                                                                                                                                                                                                                                                                                                                                                                              yongsung@mail.kribb.re.kr
2 row: D column: 09
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Hahn, Y., Oh,
Cheong, J.E.,
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                                                                                                                                                                                                                                      /cell_type="Epithelial"
/cell_line="SNU-668"
/lab_host="Top10F'"
                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="$2$NU668$1-2-D09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/tissue_type="cartilage"
/lab_host="E.coli DH10 B"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: No
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/sex="M"
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Eutheria; Primates;
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Tuseong-gu, Daejeon 305-333, South Korea
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Pred. No.
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7.9e-05;
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J.M., Park,H.S., Kim,S.
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BG538695
BG538695.1 GI
 Similarity 100 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            http://image.llnl.gov
Plate: LLCM1510 row: p column
High quality sequence stop: 499
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONTECH Laboratories, Inc.
cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
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602566932F1 NIH_MGC_77 Homo
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Mammalia;
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                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4691574"
/clone=lib="NIH_MGC_77"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
Sfil (ggccgcctcggcc); Site_2: Sfil (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A,
C, or G and N = A, C, G, or T). Average insert size 1.9
kb (range 0.5-4.0 kb). 12/15 colonies contained inserts
by PCR. This library was enriched for full-length clones
and was constructed by Clontech Laboratories (Palo Alto,
CA). Note: this is a NIH_MGC Library."
95 a 194 c 213 g 116 t
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BG928868.1 GI:14323391
EST.
human.
HOMO sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 620)
Kumar,S., Connor,J.R., Dodds,R.A., Halsey,W., Van Horn,M., Sathe,G., Mui,P., Agarwal,P., Badger,A.M., Lee,J.C., Gower Lark,M.W.
Identification and initial characterization of 5000 expressequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries
Osteoarthritic cartilage cDNA libraries
Osteoarthritic cartil. 9 (7), 641-653 (2001)
21482651
Contact: Sanjay Kumar
UW2109
GlaxOSmithKline
709 Swedeland Road, P.O. Box 1539, King of Prussia, PA 194
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mRNA sequence
BI457141
BI457141.1 G
EST.
human.
Homo sapiens
Eukaryota; Me
Mammalia; Eut
1 (bases 1 t
NIH-MGC http:
National Inst
Unpublished (
Contact: Robe
             Eukaryota; Metazoa; Chordata; Crar
Mammalia; Eutheria; Primates; Cata
1 (bases 1 to 651)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Man
Unpublished (1999)
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Fax: 610-270-5598
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03185392F1 NIH_MGC_42 Homo
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/clone_lib="HNC (Human Normal Cartilage"
/tissue_type="cartilage"
/lab_host="E.coli DH10 B"
/note="Vector: pSPORT I; Site_1:
Directional"
a 218 c 207 ~
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imer: T7.
   Robert
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1. .620
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IMAGE:5258159 5',
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Location/Qualifiers
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Plate: LLCM1869
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http://image.llnl.gov
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larity 100.0%;
Conservative
/organism="Homo sapiens"
/organism="Homo sapiens"
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/clone="CSODK007Y021"
/clone_lib="LTI_NFL006_PL2"
/fissue_type="placenta"
/note="Vector: pCMVSPORT 6; Sitell: NotI; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, adivision of Invitrogen 9800 Medical Center Drive
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/db_xref="taxon:9606"
/clone="IMAGE:5258159"
/clone="IMAGE:5258159"
/clone=lib="NIH_MGC_42"
/tissue_type="epithelioid carcinoma cell line"
/tissue_type="epithelioid carcinoma cell line"
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/lab_host="DH10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI; XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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Procurement: ATCC
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Local s
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1929 row: k column: 19
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/clone=lib="NIH_MGC_40"
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/lab_host="DH10B (phage-resistant)"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                      DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 32
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                                                                                                                                                                                                    ORGANISM
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M Homo Sapie...

Eukaryota; Metazoa; Chordata; Ciu...

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; nu.

1 (bases 1 to 800)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collectio

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

~NNA Library Preparation: Life Technologies, Inc.

(LI
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                                                                                                                                                                                                                                 BI826781
603077268F1 NIH_MGC_119 F
mRNA sequence.
BI826781
BI826781.1 GI:15938331
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.2%;
Similarity 100.0%;
32; Conservative
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Mammalia; E
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EST.
                                                                                                                                                                                                                   human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 790)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLCM1944 row: h column: 17
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BM046275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NIH_MGC_40"
/tissue_type="carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eutheria; 1 to 790)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens/db_xref="taxon:9606"/clone="IMAGE:5452480"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .790
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 32; DE Pred. No. 8. 0; Mismatches
                                                                                                                                                                                                                                                                                                   Homo
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sapiens cDNA
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Catarrhini;
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8.9e-05;
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i; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene
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                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                               Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        790;
                                                                                                                                                                                                                                                                                                   IMAGE:5169065 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Euteleostomi;
              (LLNL)
                                                                                                                                                                                Euteleostomi;
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ORIGIN
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Best Local S
Matches 32
 cch completed: Ctime: 1335.79
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                                                                                                                          Similarity 32; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be ound through the I.M.A.G.E. Consortium/LLNL at: ttp://image.llnl.gov
late: LLAM11420 row: g column: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             igh quality sequence stop:
    Location/Qualifiers
                                                              GTACCCCTGGTGCTGGATGGCTGTGGCTGCTG
                                                                                            GTACCCCTGGTGCTGGATGGCTGTGGCTGCTG 158
                                                                                                                        4.2%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                           115
July 28,
9 secs
                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5169065"
/clone_lib="NIH_MGC_119"
/tissue_type="medulla"
/lab_host="DH10B"
/note="Organ: brain; Vectorsite_2: EcoRV (destroyed)
                                                                                                                                                                                                                                                                  /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for
                2003,
                18:02:13
                                                                                                                          0;
                                                                                                                                          Score 32;
Pred. No.
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                                                                                                                        DB ls;
8.9e-05;
                                                                                                                                                                                                                                    was constructed by
Genetics tracking
                                                                                                                                                         Length 800;
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